SEQUENCE LISTING

SEQ ID NO: 1 human IPM 150 cDNA, isoform A (3330 bp)

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1 TAAACCAAGA AGGTTATCCT CAATCATCTG GTATCAATAT ATAATTATTT TTCACATTTC
  61 TGTTACTTTT TAATGAGATT TGAGGTTGTT CTGTGATTGT TATCAGAATT ACCAATGCAC
 121 AAAAGCCAGA ATGTATTTGG AAACTAGAAG AGCTATTTTT GTTTTTTGGA TTTTTCTCCA
 181 AGTTCAAGGA ACCAAAGATA TCTCCATTAA CATATACCAT TCTGAAACTA AAGACATAGA
 241 CAATCCCCCA AGAAATGAAA CAACTGAAAG TACTGAAAAA ATGTACAAAA TGTCAACTAT
 301 GAGACGAATA TTCGATTTGG CAAAGCATCG AACAAAAAGA TCCGCATTTT TCCCAACGGG
 361 GGTTAAAGTC TGTCCACAGG AATCCATGAA ACAGATTTTA GACAGTCTTC AAGCTTATTA
 421 TAGATTGAGA GTGTGTCAGG AAGCAGTATG GGAAGCATAT CGGATCTTTC TGGATCGCAT
 481 CCCTGACACA GGGGAATATC AGGACTGGGT CAGCATCTGC CAGCAGGAGA CCTTCTGCCT
 541 CTTTGACATT GGAAAAAACT TCAGCAATTC CCAGGAGCAC CTGGATCTTC TCCAGCAGAG
 601 AATAAAACAG AGAAGTTTCC CTGACAGAAA AGATGAAATA TCTGCAGAGA AGACATTGGG
 661 AGAGCCTGGT GAAACCATTG TCATTTCAAC AGCAATCTAC ATTTCAAAGA CTTGGGCAGT
 721 ATTCTAAGAA AACCCTCAGA AGAGCAAATT CAAGATGTTG CCAACGTCTC ACTTGGGCCT
 781 TTCCCTCTCA CTCCTGATGA CACCCTCCTC AATGAAATTC TCGATAATAC ACTCAACGAC
 841 ACCAAGATGC CTACAACAGA AAGAGAAACA GAATTCGCTG TGTTGGAGGA GCAGAGGGTG
 901 GAGCTCAGCG TCTCTCTGGT AAACCAGAAG TTCAAGGCAG AGCTCGCTGA CTCCCAGTCC
 961 CCATATTACC AGGAGCTAGC AGGAAAGTCC CAACTTCAGA TGCAAAAGAT ATTTAAGAAA
1021 CTTCCAGGAT TCAAAAAAAT CCATGTGTTA GGATTTAGAC CAAAGAAAGA AAAAGATGGC
1081 TCAAGCTCCA CAGAGATGCA ACTTACGGCC ATCTTTAAGA GACACAGTGC AGAAGCAAAA
1141 AGCCCTGCAA GTGACCTCCT GTCTTTTGAT TCCAACAAAA TTGAAAGTGA GGAAGTCTAT
1201 CATGGAACCA TGGAGGAGGA CAAGCAACCA GAAATCTATC TCACAGCTAC AGACCTCAAA
1261 AGGCTGATCA GCAAAGCACT AGAGGAAGAA CAATCTTTGG ATGTGGGGAC AATTCAGTTC
1321 ACTGATGAAA TTGCTGGATC ACTGCCAGCC TTTGGTCCTG ACACCCAATC AGAGCTGCCC
1381 ACATCTTTTG CTGTTATAAC AGAGGATGCT ACTTTGAGTC CAGAACTTCC
                                                            TCCTGTTGAA
1441 CCCCAGCTTG AGACAGTGGA CGGAGCAGAG CATGGTCTAC CTGACACTTC TTGGTCTCCA
1501 CCTGCTATGG CCTCTACCTC CCTGTCAGAA GCTCCACCTT TCTTTATGGC ATCAAGCATC
1561 TTCTCTCTGA CTGATCAAGG CACCACAGAT ACAATGGCCA CTGACCAGAC AATGCTAGTA
1621 CCAGGGCTCA CCATCCCCAC CAGTGATTAT TCTGCAATCA GCCAACTGGC TCTGGGAATT
1681 TCACATCCAC CTGCATCTTC AGATGACAGC CGATCAAGTG CAGGTGGCGA AGATATGGTC
1741 AGACACCTAG ATGAAATGGA TCTGTCTGAC ACTCCTGCCC CATCTGAGGT ACCAGAGCTC
1801 AGCGAATATG TTTCTGTCCC AGATCATTTC TTGGAGGATA CCACTCCTGT CTCAGCTTTA
1861 CAGTATATCA CCACTAGTTC TATGACCATT GCCCCCAAGG GCCGAGAGCT GGTAGTGTTC
1921 TTCAGTCTGC GTGTTGCTAA CATGGCCTTC TCCAACGACC TGTTCAACAA GAGCTCTCTG
1981 GAGTACCGAG CTCTGGAGCA ACAATTCACA CAGCTGCTGG TTCCATATCT ACGATCCAAT
2041 CTTACAGGAT TTAAGCAACT TGAAATACTT AACTTCAGAA ACGGGAGTGT GATTGTGAAT
2101 AGCAAAATGA AGTTTGCTAA GTCTGTGCCG TATAACCTCA CCAAGGCTGT GCACGGGGTC
2161 TTGGAGGATT TTCGTTCTGC TGCAGCCCAA CAACTCCATC TGGAAATAGA CAGCTACTCT
2221 CTCAACATTG AACCAGCTGA TCAAGCAGAT CCCTGCAAGT TCCTGGCCTG CGGCGAATTT
2281 GCCCAATGTG TAAAGAACGA ACGGACTGAG GAAGCGGAGT GTCGCTGCAA ACCAGGATAT
2341 GACAGCCAGG GGAGCCTGGA CGGTCTGGAA CCAGGCCTCT GTGGCCCTGG CACAAAGGAA
2401 TGCGAGGTCC TCCAGGGAAA GGGAGCTCCA TGCAGGTTGC CAGATCACTC TGAAAATCAA
2461 GCATACAAAA CTAGTGTTAA AAAGTTCCAA AATCAACAAA ATAACAAGGT AATCAGTAAA
2521 AGAAATTCTG AATTACTGAC CGTAGAATAT GAAGAATTTA ACCATCAAGA TTGGGAAGGA
2581 AATTAAAAAC TGAAAATGTA CAATTATCAC TTAGGCTATC TCAAGAGAGA TGATTTGCCT
2641 TCTCAAGGAA AATGGAGACA GGCATATTCA TGGGTCATCA AAATCCAGAC ATACAGTCAA
2701 CACTGAGAAT CAGCACACAC CATATTTCAA ATATAGAAGA GTCATGTACT TGGCAACCAG
2761 TAAATTCTGA AAAAAAAGAC ACTTACTTAT TATTAAAACC CCAAATGCAA TCAGCGAAAC
2821 ATATTTTTAC TATTCTTGGA TGATAGTCAA AATGATCATA AGCCAGGTTT GCTTCCACCT
2881 TCCCTGAAAA TTTTACTCAC AGATCATTTG CAACAAGCAT AGCTTACTTA TTGTTTAGGG
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2941 ACTGAACAAT TTATTGGGAA GCAAACTCTT TATATGCTAG AAAGTACATT TAAAAGATGA
3001 CTACTTACGC AGGGAGATGC AGGTCTCTCT AAACGCATGA ATGTATGTAG TGTGTAGGCA
3061 CTGTAGTGAG TGTATATATG CTCCACACTA CGTCTGATAA ACACAAACCT CAGTATTCAG
3121 TTATTAGGCA CACTAGTTTT ATACGCAACT ACTGCTTACA TAGTAGACTG TTTTGTTGCC
3181 AATAATCTTT GAATTGTTCT TTAAAAGAAA CTGAGGTTCA GATACACATA CCATGGAAAA
3241 ATCTTACTTT TCTTGTTACT ACACAAAGCT ATTTTAAAGA AGATGCTATG TTGGGAGAAG
3301 GGCGAAGTTG TACTATATGA CATAATCAAT
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SEQ ID NO: 2 Human IPM 150 amino acid sequence, isoform A

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1 MYLETRRAIF VFWIFLQVQG TKDISINIYH SETKDIDNPP RNETTESTEK MYKMSTMRRI
61 FDLAKHRTKR SAFFPTGVKV CPQESMKQIL DSLQAYYRLR VCQEAVWEAY RIFLDRIPDT
121 GEYQDWVSIC QQETFCLFDI GKNFSNSQEH LDLLQQRIKQ RSFPDRKDEI SAEKTLGEPG
181 ETIVISTDVA NVSLGPFPLT PDDTLLNEIL DNTLNDTKMP TTERETEFAV LEEQRVELSV
241 SLVNQKFKAE LADSQSPYYQ ELAGKSQLQM QKIFKKLPGF KKIHVLGFRP KKEKDGSSST
301 EMQLTAIFKR HSAEAKSPAS DLLSFDSNKI ESEEVYHGTM EEDKQPEIYL TATDLKRLIS
361 KALEEEQSLD VGTIQFTDEI AGSLPAFGPD TQSELPTSFA VITEDATLSP ELPPVEPQLE
421 TVDGAEHGLP DTSWSPPAMA STSLSEAPPF FMASSIFSLT DQGTTDTMAT DQTMLVPGLT
481 IPTSDYSAIS QLALGISHPP ASSDDSRSSA GGEDMVRHLD EMDLSDTPAP SEVPELSEYV
541 SVPDHFLEDT TPVSALQYIT TSSMTIAPKG RELVVFFSLR VANMAFSNDL FNKSSLEYRA
601 LEQQFTQLLV PYLRSNLTGF KQLEILNFRN GSVIVNSKMK FAKSVPYNLT KAVHGVLEDF
661 RSAAAQQLHL EIDSYSLNIE PADQADPCKF LACGEFAQCV KNERTEEAEC RCKPGYDSQG
721 SLDGLEPGLC GPGTKECEVL QGKGAPCRLP DHSENQAYKT SVKKFQNQQN NKVISKRNSE
781 LLTVEYEEFN HQDWEGN
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SEQ ID NO: 3 Human IPM 150 cDNA sequence, isoform B

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1 GGGAGCTATT TTTGTTTTTT GGATTTTTCT CCAAGTTCAA GGAACCAAAG TGTGTCAGGA
  61 AGCAGTATGG GAAGCATATC GGATCTTTCT GGATCGCATC CCTGACACAG GGGAATATCA
 121 GGACTGGGTC AGCATCTGCC AGCAGGAGAC CTTCTGCCTC TTTGACATTG GAAAAAACTT
 181 CAGCAATTCC CAGGAGCACC TGGATCTTCT CCAGCAGAGA ATAAAACAGA GAAGTTTCCC
 241 TGACAGAAAA GATGAAATAT CTGCAGAGAA GACATTGGGA GAGCCTGGTG AAACCATTGT
 301 CATTTCAACA GATGTTGCCA ACGTCTCACT TGGGCCTTTC CCTCTCACTC CTGATGACAC
 361 CCTCCTCAAT GAAATTCTCG ATAATACACT CAACGACACC AAGATGCCTA CAACAGAAAG
 421 AGAAACAGAA TTCGCTGTGT TGGAGGAGCA GAGGGTGGAG CTCAGCGTCT CTCTGGTAAA
 481 CCAGAAGTTC AAGGCAGAGC TCGCTGACTC CCAGTCCCCA TATTACCAGG AGCTAGCAGG
 541 AAAGTCCCAA CTTCAGATGC AAAAGATATT TAAGAAACTT CCAGGATTCA AAAAAATCCA
 601 TGTGTTAGGA TTTAGACCAA AGAAAGAAAA AGATGGCTCA AGCTCCACAG AGATGCAACT
 661 TACGGCCATC TTTAAGAGAC ACAGTGCAGA AGCAAAAAGC CCTGCAAGTG ACCTCCTGTC
 721 TTTTGATTCC AACAAATTG AAAGTGAGGA AGTCTATCAT GGAACCATGG AGGAGGACAA
 781 GCAACCAGAA ATCTATCTCA CAGCTACAGA CCTCAAAAGG CTGATCAGCA AAGCACTAGA
 841 GGAAGAACAA TCTTTGGATG TGGGGACAAT TCAGTTCACT GATGAAATTG CTGGATCACT
 901 GCCAGCCTTT GGTCCTGACA CCCAATCAGA GCTGCCCACA TCTTTTGCTG TTATAACAGA
 961 GGATGCTACT TTGAGTCCAG AACTTCCTCC TGTTGAACCC CAGCTTGAGA CAGTGGACGG
1021 AGCAGAGCAT GGTCTACCTG ACACTTCTTG GTCTCCACCT GCTATGGCCT CTACCTCCCT
1081 GTCAGAAGCT CCACCTTTCT TTATGGCATC AAGCATCTTC TCTCTGACTG ATCAAGGCAC
1141 CACAGATACA ATGGCCACTG ACCAGACAAT GCTAGTACCA GGGCTCACCA TCCCCACCAG
1201 TGATTATTCT GCAATCAGCC AACTGGCTCT GGGAATTTCA CATCCACCTG CATCTTCAGA
1261 TGACAGCCGA TCAAGTGCAG GTGGCGAAGA TATGGTCAGA CACCTAGATG AAATGGATCT
1321 GTCTGACACT CCTGCCCCAT CTGAGGTACC AGGGCTCAGC GAATACGTTT CTGTCCCAGA
1381 TCATTTCTTG GAGGATACCA CTCCTGTCTC AGCTTTACAG TATATCACCA CTAGTTCTAT
1441 GACCATTGCC CCCAAGGGCC GAGAGCTGGT AGTGTTCTTC AGTCTGCGTG TTGCTAACAT
1501 GGCCTTCTCC AACGACCTGT TCAACAAGAG CTCTCTGGAG TACCGAGCTC TGGAGCAACA
1561 ATTCACACAG CTGCTGGTTC CATATCTACG ATCCAATCTT ACAGGATTTA AGCAACTTGA
1621 AATACTTAAC TTCAGAAACG GGAGTGTGAT TGTGAATAGC AAAATGAAGT TTGCTAAGTC
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1681 TGTGCCGTAT AACCTCACCA AGGCTGTGCA CGGGGTCTTG GAGGATTTTC GTTCTGCTGC 1741 AGCCCAACAA CTCCATCTGG AAATAGACAG CTACTCTCTC AACATTGAAC CAGCTGATCA 1801 AGCAGATCCC TGCAAGTTCC TGGCCTGCGG CGAATTTGCC CAATGTGTAA AGAACGAACG 1861 GACTGAGGAA GCGGAGTGTC GCTGCAAACC AGGATATGAC AGCCAGGGGA GCCTGGACGG 1921 TCTGGAACCA GGCCTCTGTG GCCCTGGCAC AAAGGAATGC GAGGTCCTCC AGGGAAAGGG 1981 AGCTCCATGC AGGTTGCCAG ATCACTCTGA AAATCAAGCA TACAAAACTA GTGTTAAAAA 2041 GTTCCAAAAT CAACAAAATA ACAAGGTAAT CAGTAAAAGA AATTCTGAAT TACTGACCGT 2101 AGAATATGAA GAATTTAACC ATCAAGATTG GGAAGGAAAT TAAAAACTGA AAATGTACAA 2161 TTATCACTTA GGCTATCTCA AGAGAGATGA TTTGCCTTCT CAAGGAAAAT GGAGACAGGC 2221 ATATTCATGG GTCATCAAAA TCCAGACATA CAGTCAACAC TGAGAATCAG CACACCAT 2281 ATTTCAAATA TAGAAGAGTC ATGTACTTGG CAACCAGTAA ATTCTGAAAA AAAAGACACT 2341 TACTTATTAT TAAAACCCCA AATGCAATCA GCGAAACATA TTTTTACTAT TCTTGGATGA 2401 TAGTCAAAAT GATCATAAGC CAGGTTTGCT TCCACCTTCC CTGAAAATTT TACTCACAGA 2461 TCATTTGCAA CAAGCATAGC TTACTTATTG TTTAGGGACT GAACAATTTA TTGGGAAGCA 2521 AACTCTTTAT ATGCTAGAAA GTACATTTAA AAGATGACTA CTTACGCAGG GAGATGCAGG 2581 TCTCTCTAAA CGCATGAATG TATGTAGTGT GTAGGCACTG TAGTGAGTGT ATATATGCTC 2641 CACACTACGT CTGATAAACA CAAACCTCAG TATTCAGTTA TTAGGCACAC TAGTTTTATA 2701 CGCAACTACT GCTTACATAG TAGACTGTTT TGTTGCCAAT AATCTTTGAA TTGTTCTTTA 2761 AAAGAAACTG AGGTTCAGAT ACACATACCA TGGAAAAATC TTACTTTTCT TGTTACTACA 2821 CAAAGCTATT TTAAAGAAGA TGCTATGTTG GGAGAAGGGC GAAGTTGTAC TATATGACAT 2881 AATCAAT

SEQ ID NO: 4 Human IPM 150 amino acid sequence, isoform B

1 MYLETRRAIF VFWIFLQVQG TKVCQEAVWE AYRIFLDRIP DTGEYQDWVS ICQQETFCLF
61 DIGKNFSNSQ EHLDLLQQRI KQRSFPDRKD EISAEKTLGE PGETIVISTD VANVSLGPFP
121 LTPDDTLLNE ILDNTLNDTK MPTTERETEF AVLEEQRVEL SVSLVNQKFK AELADSQSPY
181 YQELAGKSQL QMQKIFKKLP GFKKIHVLGF RPKKEKDGSS STEMQLTAIF KRHSAEAKSP
241 ASDLLSFDSN KIESEEVYHG TMEEDKQPEI YLTATDLKRL ISKALEEEQS LDVGTIQFTD
301 EIAGSLPAFG PDTQSELPTS FAVITEDATL SPELPPVEPQ LETVDGAEHG LPDTSWSPPA
361 MASTSLSEAP PFFMASSIFS LTDQGTTDTM ATDQTMLVPG LTIPTSDYSA ISQLALGISH
421 PPASSDDSRS SAGGEDMVRH LDEMDLSDTP APSEVPGLSE YVSVPDHFLE DTTPVSALQY
481 ITTSSMTIAP KGRELVVFFS LRVANMAFSN DLFNKSSLEY RALEQQFTQL LVPYLRSNLT
541 GFKQLEILNF RNGSVIVNSK MKFAKSVPYN LTKAVHGVLE DFRSAAAQQL HLEIDSYSLN
601 IEPADQADPC KFLACGEFAQ CVKNERTEEA ECRCKPGYDS QGSLDGLEPG LCGPGTKECE
661 VLQGKGAPCR LPDHSENQAY KTSVKKFQNQ QNNKVISKRN SELLTVEYEE FNHQDWEGN

SEQ ID NO: 5 Human IPM 150 cDNA sequence, isoform C

1 AAATTAACAC CCTCATAAAG GTAAACCAAG AAGGTTATCC TCAATCATCT GGTATCAATA
61 TATAATTATT TTTCACATTT CTGTTACTTT TTAATGAGAT TTGAGGTTGT CTGTGATTGT
121 TATCAGAATT ACCAATGCAC AAAAGCCAGA ATGTATTTGG AAACTAGAAG AGCTATTTTT
181 GTTTTTTGGA TTTTTCTCCA AGTTCAAGGA ACCAAAGATA TCTCCATTAA CATATACCAT
241 TCTGAAACTA AAGACATAGA CAATNCCCCA AGAAATGAAA CAACTGAAAG TACTGAAAAA
301 ATGTACAAAA TGTCAACTAT GAGACGAATA TTCGATTTGG CAAAGNATCG AACAAAAAGA
361 TCCGCATTTT TCCCAACGGG GGTTAAAGTC TGTCCACAGG AATCCATGAA ACAGATTTTA
421 GACAGTCTTC AAGCTTATTA TAGATTGAGA GTGTGTCAGG AAGCAGCATG GGAAGCATAT
481 CGGATCTTC TGGATCGCAT CCCTGACACA GGGGAATATC AGGACTGGGT CAGCATCTGC
541 CAGCAGGAGA CCTTCTGCCT CTTTGACATT GGAAAAAACT TCAGCAATTC CCAGGAGCAC
601 CTGGATCTTC TCCAGCAGAG AATAAAACAG AGAAGTTTCC CTGACAGAAA AGATGAAATA
661 TCTGCAGAGA AGACATTGGG AGACCCTGGT GAAACCATTG TCATTTCAAC AGCAATCTAC
721 ATTTCAAAGA CTTGGGCCT TTCCCTCTCA CTCCTGATGA CACCCTCCTC AATGGAATT
841 TCGATAATAC ACTCAACGAC ACCAAGATGC CTACAACAGA AAGAGAAACA GAATTCGCTG
901 TGTTGGAGGA GCAGAGGGTG GAGCTCAGCG TCTCTCTGGT AAACCAGAAG TTCAAGGCAG

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961 AGCTCGCTGA CTCCCAGTCC CCATATTACC AGGAGCTAGC AGGAAAGTCC CAACTTCAGA
1021 TGCAAAAGAT ATTTAAGAAA CTTCCAGGAT TCAAAAAAAT CCATGTGTTA GGATTTAGAC
1081 CAAAGAAGA AAAAGATGGC TCAAGCTCCA CAGAGATGCA ACTTACGGCC ATCTTTAAGA
1141 GACACAGTGC AGAAGCAAAA AGCCCTGCAA GTGACCTCCT GTCTTTTGAT TCCAACAAAA
1201 TTGAAAGTGA GGAAGTCTAT CATGGAACCA TGGAGGAGGA CAAGCAACCA GAAATCTATC
1261 TCACAGCTAC AGACCTCAAA AGGCTGATCA GCAAAGCACT AGAGGAAGAA CAATCTTTGG
1321 ATGTGGGGAC AATTCAGTTC ACTGATGAAA TTGCTGGATC ACTGCCAGCC TTTGGTCCTG
1381 ACACCCAATC AGAGCTGCCC ACATCTTTTG CTGTTATAAC AGAGGATGCT ACTTTGAGTC
1441 CAGAACTTCC TCCTGTTGAA CCCCAGCTTG AGACAGTGGA CGGAGCAGAG CATGGTCTAC
1501 CTGACACTTC TTGGTCTCCA CCTGCTATGG CCCTACCTCC CTGTCAGAAG CTCCACCTTT
1561 CTTTATGGCA TCAAGCATCT TCTCTCTGAC TGATCAAGGC ACCACAGATA CAATGGCCAC
1621 TGACCAGACA ATGCTAGTAC CAGGGCTCAC CATCCCCACC AGTGATTATT CTGCAATCAG
1681 CCAACTGGCT CTGGGAATTT CACATCCACC TGCATCTTCA GATGACAGCC GATCAAGTGC
1741 AGGTGGCGAA GGTATGGACA GAGACCTAGA TGAAATGGAT CTGTCTGACA CTCCTGCCCC
1801 ATCTGAGGTA CCAGAGCTCA GCGAATATGT TTCTGTCCCA GATCATTTCT TGGAGGATAC
1861 CACTCCTGTC TCAGCTTTAC AGTATATCAC CACTAGTTCT ATGACCATTG CCCCCAAGGG
1921 CCGAGAGCTG GTAGTGTTCT TCAGTCTGCG TGTTGCTAAC ATGGCCTTCT CCAACGACCT
1981 GTTCAACAAG AGCTATTTGG AGTACCGAGC TCTGGAGCAA CAATTCACAC AGCTGCTGGT
2041 TCCATATCTA CGATCCAATC TTACAGGATT TAAGCAACTT GAAATACTTA ACTTCAGAAA
2101 CGGGAGTGTG ATTGTGAATA GCAAAATGAA GTTTGCTAAG TCAGTGCCGT ATAACCTCAC
2161 CAAGGCTGTG CACGGGGTCT TGGAGGATTT TCGTTCTGCT GCAGCCCAAC AACTCCATCT
2221 GGAAATAGAC AGCTACTCTC TCCC
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SEQ ID NO: 6 Human IPM 150 amino acid sequence, isoform C

1 MYLETRRAIF VFWIFLQVQG TKDISINIYH SETKDIDNPP RNETTESTEK MYKMSTMRRI 61 FDLAKHRTKR SAFFPTGVKV CPQESMKQIL DSLQAYYRLR VCQEAAWEAY RIFLDRIPDT 121 GEYQDWVSIC QQETFCLFDI GKNFSNSQEH LDLLQQRIKQ RSFPDRKDEI SAEKTLGEPG 181 ETIVISTAIY ISKTWAVF

SEQ ID NO: 7 Human IMPG1 gene, regulatory region

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1 AGGGTGTAGG CTTTTGAACC AGGACTCTTT AGGTTTAAAT CCTAGCTCTG CCACATATAC
  61 TTTATTCTCC TCAAATTTAA AAGAGATAGT ATTAACAGTG TTTATATTGT CATATTGAGG
 121 AATCTATGGA TAATCTATGG ACATCTCTAA GAACAATGTC TATCCACAAC ACAAGAGCTC
 181 AATATACAGT AGTAGTTGCA GTGTGTTTCA TGACTCAGCA ATATGTAGCA TGTATAGTCA
 241 AAATAATATA AAATCAAATA TTCAAAAACT GAAATTACAA TAATACTGAT GAAGAAAGAT
 301 GGAAAGATGT TTACAATGAG TAGAAAGGGT ATGTGTGGAA GTGAAGTTAT TCTCAATATC
 361 TATTATTTGA TAATACCTAA AAGTGAAAAC CTCCAAAATA GTAATAGAGG CATGTTATTT
 421 AGAAGTGCAA ATGAGACTAC TAGAAGAATT AGGTTGATGA AGTAAAAATG GCTCCCCTTT
 481 GAAAGAAGGC ATGGGTAGAA GAAAGGCACA ATTTTTTCTT ACAAACTTTG TAGAAAAAAA
 541 GTATTTGACC CCTTAAACAC AGTGCATACA GATTTTAAAC ATTAAAACCA GACTTAAATC
 601 AAAAAAGCCA CCTGTATGTA ATTCCAAATC AAAAGCAATT TATAAAGCAG AACATAGAAG
 661 AGAATGGAGA CAGTTTCGCT ATCTGTGGAG ACTAATACAT ATTGGATAAC CATATACTTT
721 CAGGGACAGA AATTAAGCTC TTTTAATGGA TGTTTCTTGT ACATGTCATT TTAGAAAACA
781 TCTGACCCTA ACTGTCAGCC TTATTCTCTG TTTGGCAGAA CTTCCCCTGG CTCTCTGTGT
841 CACTGTAACA GGTGAATAAC TAAGAAAAAA CTGTGTCTGT AGACACTTGT TTATAATGGC
 901 ATTCAGGGTC CTGGAGCTAG GCTGACAGAT GCTCCTCCAG AAGGTTAATG AGATAAAGGT
 961 TCCTCCAGCT GGCCCTTAAG CAGAGATTAC ACCTGAGGGA AAGACAAGCA GATTATTCCA
1021 GAAACAGACA CTGCTACATG TTCTTCATAA ATTAACACCC TCATAAAGGT AAACCAAGAA
1081 GGTTATCCTC AATCATCTGG TATCAATATA TAATTATTTT TCACATTTCT GTTACTTTTT
1141 AATGAGATTT GAGGTTGTTC TGTGATTGTT ATCAGAATTA CCAATGCACA AAAGCCAGAA
1201 TGTATTTGGA AACTAGAAGA GCTATTTTTG TTTTTTTGGAT TTTTCTCCAA GTTCAAGGAA
1261 CCAAAGGTAA GTTACTTAAA TGTTTACTTT TAAATTGCTT ATCTATAAAA TCTACCGATA
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1321 GAAGTGAATA TTTAGAACCA ACAAGGCTAC CAATTTATCT CACGGGCTAG TATATAGTAG
1381 GCCTTGAATA AATATTGCTT GATTGATTGA ATAATTAACT ATCAGAAATG ATTTTCACTT
1441 GATTTAATAT TTACTACATG GTCTTAAGTG CAGTGAAGAT TAACAAAATA GGAGAGATGA
1501 ATGCATCCTA TTTGCTGTTC TAAAACATTC ATTGAAAATT CTTATTATTA AATGTAAATA
1561 NTATTAGTAG ATCTGGTGAA AACTAAACTC CATTTATCCA CCCGAAATTC AACCAAATAA
1621 AACCTAAAGG ATAAAAGTAA TGTTTTAAGT CATTTATGGT CAGACAAAAA AAAGTAAGTA
1681 TTTCTTACCT TCTCACAATG AAATCATGAG TTGCTTTCCC TTAGAAAATA GCAAACATTC
1741 TTCATCTTCA GGGTTCATGA TGACAACCAC TTCAAAATTT GGTTGTTTTT GAAAGTTGTA
1801 CGCATAAAAG AACTAGGCAA TGTATGTTCT TATGGCAAAT CTGCATCTGA ATATGAAA

SEQ ID NO: 8 Mouse IPM 150 cDNA sequence, isoform A

1 ACAAGATTAT TCCAGGAACT GACCTGCTTC CGGATCCTCG AGAATTAGCA CCTTCATAGG 61 GTAAACCAGG GAGGTCGTCT GTACTCAGCC GGCACCTGGA TTTGATTATT TTTCATATTT 121 CAGTCACCTT ATTTCTTTAA GTGTGACTTG GTATTGTTCT GTGATTTTTC AGAATTACCA 181 GTATACAGAA CCAGAATGAA TTTTCAAATT AAACATGCTA TCTTTGTTTT TGGGATTTTT 241 CTCCAAGTTC AAGGAATCAA AGATACCTCT ATTAAAATAT TCAGTTCTGA AATTAAAAAC 301 ATAGACAAAA CCCCAAGAAT CGAAACAATT GAAAGTACTT CAACAGTGCA CAAAGTGTCA 361 ACCATGAAAC GAATATTCGA TTTGCCAAAG CTTCGAACCA AAAGATCAGC ACTTTTCCCA 421 GCTGCTAACA TCTGTCCACA GGAATCCTTG AGACAGATTT TAGCAAGTCT TCAAGAATAT 481 TATAGACTGA GAGTATGTCA AGAAGTCGTG TGGGAAGCAT ATCGTATCTT TCTGGACCGA 541 ATTCCTGACA CAGAGGAATA TCAAGACTGG GTCAGCCTCT GCCAGAAAGA AACCTTCTGC 601 CTCTTTGACA TTGGGAAAAA CTTCAGCAAC TCCCAGGAGC ACCTAGATCT TCTTCAGCAG 661 AGAATAAAAC AGAGAAGCTT CCCTGGGAGG AAAGATGAGA CAGCCTCCAT GGAGACACTG 721 GAAGCACCTA CTGAAGCCCC TGTGGTACCC ACAGATGTTT CCAGGATGTC CCTGGGGCCC 781 TTCCCACTTC CTTCTGATGA CACAGACCTC AAGGAGATTC TCAGTGTCAC CCTCAAGGAC 841 ATTCAAAAGC CCACAACAGA AAGTAAAACA GAACCTATTC ACGTGTCTGA ATTCTCATCA 901 GAGGAGAAGG TGGAATTCAG CATCTCTCTG CCAAACCACA GGTTCAAGGC AGAGCTCACC 961 AACTCTGGGT CACCATACTA CCAGGAACTG GTGGGACAGT CCCAACTGCA GTTGCAAAAG 1021 ATATTTAAGA AACTTCCAGG ATTCGGAGAA ATCCGTGTAT TAGGATTTAG ACCAAAGAAA 1081 GAAGAAGATG GTTCAAGCTC CACAGAAATA CAGCTTATGG CCATCTTTAA GAGGGACCAT 1141 GCAGAAGCAA AAAGCCCTGA TAGTCATCTA CTGTCTCTTG ATTCCAACAA AATTGAAAGT 1201 GAAAGAATCC ATCATGGAGT CATAGAAGAC AAACAACCAG AAACCTACCT CACAGCTACA 1261 GACCTCAAAA AACTCATCAT ACAACTACTA GATGGAGACC TGTCCTTGGT AGAAGGGAAA 1321 ATTCCATTCG GTGATGAAGT TACTGGGACA CTCTTCAGAC CTGTCACTGA ACCAGATCTG 1381 CCCAAGCCCC TTGCTGATGT CACAGAGGAT GCCACTTTGA GTCCAGAACT TCCTTTCGTT 1441 GAGCCTAGGC TTGAGGCAGT GGACAGAGAA GGATCTGAGC TGCCTGGAAT GTCCTCCAAA 1501 GACAGTTCTT GGTCTCCACC TGTATCAGCC TCAATTTCCC GATCAGAAAA TCTACCTTCG 1561 TTTACACCTA GCATCTTCTC TCTAGATGCT CAAAGCCCCC CTCCCTTGAT GACCACTGGC 1621 CCAACAGCAC TCATCCCCAA GCCCACTCTC CCCACTATCG ATTATTCTAC CATCCGCCAA 1681 TTGCCTCTGG AATCGTCACA TTGGCCTGCA TCCTCCAGTG ACAGAGAGCT GATCACAAGC 1741 AGCCATGACA CAATCCGAGA CCTAGATGGC ATGGATGTGT CTGACACGCC AGCCTTGTCA 1801 GAAATATCAG AACTGAGTGG ATACGATTCT GCCTCGGGTC AGTTCTTGGA GATGACCACA 1861 CCCATCCCAA CAGTACGGTT CATCACCACC AGCTCCGAGA CCATTGCCAC CAAGGGCCAG 1921 GAGCTAGTGG TATTCTTCAG CCTGCGTGTT GCTAACATGC CGTTCTCCTA TGACCTGTTC 1981 AACAAGAGTT CTCTGGAGTA TCAAGCCCTG GAACAACGAT TCACAGACCT GCTGGTTCCC 2041 TATCTACGAT CGAATCTTAC GGGATTTAAG CAACTGGAAA TACTCAGCTT CAGAAACGGA 2101 AGTGTGATCG TGAACAGCAA AGTGCGGTTT GCAAAGGCGG TACCCTACAA CCTCACCCAG 2161 GCCGTGCGCG GGGTCTTGGA GGATCTTCGG TCCACCGCAG CTCAAGGGCT CAATCTGGAA 2221 ATCGAAAGCT ACTCCCTCGA CATTGAACCA GCTGATCAGG CGGATCCCTG CAAACTCCTA 2281 GACTGTGGCA AATTTGCCCA GTGTGTAAAG AATGAGTGGA CAGAGGAAGC AGAGTGTCGC 2341 TGCAGACAGG GACATGAGAG CCACGGGACC CTGGACTACC AGACCCTGAA CCTCTGTCCC 2401 CCTGGAAAGA CTTGTGTGGC CGGCCGAGAA CAAGCAACTC CATGCAGGCC ACCAGATCAC 2461 TCTACAAACC AAGCTCAGGA ACCTGGTGTT AAAAAGCTAC GTCAGCAAAA TAAGGTAGTC 2521 AAGAAAAGAA ATTCTAAACT ATCAGCTATA GGATTTGAAG AATTTGAAGA CCAGGACTGG 2581 GAGGGAAATT AAAAGCTGGA ATCATATGCA TTATGTTGCA AACTCTGTTG AAAGGAAACT

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2641 TTATTTCTTA AAGAAAGGTG TATCTGTTCT GTTAACTTCT GAAAAACAGA GGGAGAGATT
2701 CAGTGGTCAT TGGAATACAG GCATGTAATC AACTTTGAGA CTCAGCATGC TTGAACAAGA
2761 GCACAGGCGT GTATTTGATG ACAGTTAAGC CTGGTTGGGG CGGGGGGCAC ATATTTTTAG
2821 TCAAAACTCA AAGCAATCAT TGGAACACAT TTGACTATTT TTGGACAGTA CTCAAGTAGC
2881 AAAGATAAGG TTAGCTTTTT TCTTTCTTTA AATTATTACA TAAARCTTAT TTCAAATAAA
2941 TACAACTTGT TTAGTGGGTT GTACAATATT GAGGATCTGA TTCTTTTATA TGTTAGAATA
3001 TACAGTTAAA AGATTATCAT TTGGGCCAGA GAGATAGCTA AGTGGTTAAG AGTATATACT
3061 GCTCTTCCAG AAGCCCTGGG TTTACCGTCC CAACAGCCAC ATTGACTGGC TCACACACAC
3121 CTGTAAGTCA GGCTCCAGAG AACAAACACC CTCCTCTGGC CTTTGTACCC ACGTGCACAT
3181 AACCGCAAAC AGACACACCC ACGCTATTTT TTTAGAAGTC ATTGATTTTT TTAATTAGGG
3241 GTGGAAAAKC AGGCTGGAGA GATGACTCCG TGGTTAAGAA CAGTTGTTGT TCTTCCAGAG
3301 GACCCAGGTT CAGTTCCCAG AACCCACAAG GCNAGTCTCC CAACTATTCA TAATTCTAGT
3361 TCAAGTGGAT CCAGCACCCT CTTCTAACTG ATACTGCCAG TACCAGGCAG CCATGTGGTG
3421 CATATGCATT TGGGCAGGTA AAACACTCAG ACACGCAAAA AATTTTAAAT CTAAATTTTG
3481 AAAATATTTT AGTTTTAAGG ATGATCACTG TGTGAGGGTC AGGTCTCTTA TGTATGAATG
3541 TAGTACCAAG AACTGTGATG AGTATATGTA TGCTCCATTC TATAGTCTCC TCTCTCTCTC
3601 TCTCTCTCTC TCTCTCTCT TCTCTCTCT TCTCTGGAAT TCCGGAATTC CGGAATTCCG
3661 GAATTCCG
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SEQ ID NO: 9 Mouse IPM 150 amino acid sequence, isoform A

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1 MNFQIKHAIF VFGIFLQVQG IKDTSIKIFS SEIKNIDKTP RIETIESTST VHKVSTMKRI
61 FDLPKLRTKR SALFPAANIC PQESLRQILA SLQEYYRLRV CQEVVWEAYR IFLDRIPDTE
121 EYQDWVSLCQ KETFCLFDIG KNFSNSQEHL DLLQQRIKQR SFPGRKDETA SMETLEAPTE
181 APVVPTDVSR MSLGPFPLPS DDTDLKEILS VTLKDIQKPT TESKTEPIHV SEFSSEEKVE
241 FSISLPNHRF KAELTNSGSP YYQELVGQSQ LQLQKIFKKL PGFGEIRVLG FRPKKEEDGS
301 SSTEIQLMAI FKRDHAEAKS PDSHLLSLDS NKIESERIHH GVIEDKQPET YLTATDLKKL
361 IIQLLDGDLS LVEGKIPFGD EVTGTLFRPV TEPDLPKPLA DVTEDATLSP ELPFVEPRLE
421 AVDREGSELP GMSSKDSSWS PPVSASISRS ENLPSFTPSI FSLDAQSPPP LMTTGPTALI
481 PKPTLPTIDY STIRQLPLES SHWPASSSDR ELITSSHDTI RDLDGMDVSD TPALSEISEL
541 SGYDSASGQF LEMTTPIPTV RFITTSSETI ATKGQELVVF FSLRVANMPF SYDLFNKSSL
601 EYQALEQRFT DLLVPYLRSN LTGFKQLEIL SFRNGSVIVN SKVRFAKAVP YNLTQAVRGV
661 LEDLRSTAAQ GLNLEIESYS LDIEPADQAD PCKLLDCGKF AQCVKNEWTE EAECRCRQGH
721 ESHGTLDYQT LNLCPPGKTC VAGREQATPC RPPDHSTNQA QEPGVKKLRQ QNKVVKKRNS
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SEQ ID NO:10 Mouse IPM 150 cDNA sequence, isoform D

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1 GGCTTTAAAC CAGGGAGGTC GTCTGTACTC AGCCGGCACC TGGATTTGAT TATTTTCAT 61 ATTTCAGTCA CCTTATTTCT TTAAGTGTGA CTTGGTATTG TTCTGTGATT TTTCAGAATT 121 ACCAGTATAC AGAACCAGAA TGAATTTTCA AATTAAACAT GCTATCTTTG TTTTTGGGAT 181 TTTCTCCAA GTTCAAGGAA TCAAAGTATG TCAAGAAGTC GTGTGGGAAG CATATCGTAT 241 CTTTCTGGAC CGAATTCCTG ACACAGAGGA ATATCAAGAC TGGGTCAGCC TCTGCCAGAA 301 AGAAACCTTC TGCCTCTTTG ACATTGGGAA AAACTTCAGC AACTCCCAGG AGCACCTAGA 361 TCTTCTTCAG CAGAGAATAA AACAGAGAAG CTTCCCTGGG AGGAAAGATG AGACAGCCTC 421 CATGGAGACA CTGGAAGCAC CTACTGAAGC CCCTTGGGTA CCCACAGATG TTTCCAGGAT 481 GTCCCTGGGG CCATTCCCAC TTCCTTCTGA TGACACAGAC CTCAAGGAGA TTCTCAGTGT 541 CACCCTCAAG GACATTCAAA AGCCCACAAC AGAAAGTATA ACAGAACCTA TTCACGTGTC 601 TGAATTCTCA TCAGAGGAGA AGGTGGAGTT CAGCATCTCT CTGCCAAACC ACAGGTTCAA 661 GGCAGAGCTC ACCAACTCTG GGTCACCATA CTACCAGGAA CTGGTGGGAC AGTCCCAACT 721 GCAGTTGCAA AAGATATTTA AGAAACTTC AGGATTCGGA GAAATCCGTG TATTAGGATT 781 TAGACCAAAG AAAGAAGAAG CAAAAAGCCC TGATAGTCAT CTACCTGTCT TTGATTCCAA 901 CAAAAATTGAA AGTGAAAGAA TCCATCATG AGTCATAGAA GACAAACAAC CAGAAACCTA
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SEQ ID NO:11 Mouse IPM 150 amino acid sequence, isoform D

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1 MNFQIKHAIF VFGIFLQVQG IKVCQEVVWE AYRIFLDRIP DTEEYQDWVS LCQKETFCLF
61 DIGKNFSNSQ EHLDLLQQRI KQRSFPGRKD ETASMETLEA PTEAPVVPTD VSRMSLGPFP
121 LPSDDTDLKE ILSVTLKDIQ KPTTESITEP IHVSEFSSEE KVEFSISLPN HRFKAELTNS
181 GSPYYQELVG QSQLQLQKIF KKLPGFGEIR VLGFRPKKEE DGSSSTEIQL MAIFKRDHAE
241 AKSPDSHLLS LDSNKIESER IHHGVIEDKQ PETYLTATDL KKLIIQLLDG DLSLVEGKIP
301 FGDEVTGTLF RPVTEPDLPK PLADVTEDAT LSPELPFVEP RLEAVDREGS ELPADQADPC
361 KLLDCGKFAQ CVKNEWTEEA ECRCRQGHES HGTLDYQTLN LCPPGKTCVA GREQATPCRP
TDHSTNQAQE PGVKKLRQQN KVVKKRNSKL SAIGFEEFED QDWEGN
//
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SEQ ID NO:12 Mouse IPM 150 cDNA sequence, isoform E

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1 TCTGCAGAAT TCGGCTTAAC CAGGGAGGTC GTCTGTACTC AGCCGGCACC TGGATTTGAT
  61 TATTTTTCAT ATTTCAGTCA CCTTATTTCT TTAAGTGTGA CTTGGTATTG TTCTGTGATT
 121 TTTCAGAATT ACCAGTATAC AGAACCAGAA TGAATTTTCA AATTAAACAT GCTATCTTTG
 181 TTTTTGGGAT TTTTCTCCAA GTTCAAGGAA TCAAAGATAC CTCTATTAAA ATATTCAGTT
 241 CTGAAATTAA AAACATAGAC AAAACCCCAA GAATCGAAAC AATTGAAAGT ACTTCAACAG
 301 TGCACAAAGT GTCAACCATG AAACGCCAGC CTTGTCAGAA ATATCAGAAC TGAGTGGATA
 361 CGATTCTGCC TCGGGTCAGT TCTTGGAGAT GACCACACCC ATCCCAACAG TACGGTTCAT
 421 CACCACCAGC TCCGAGACCA TTGCCACCAA GGGCCAGGAG CTAGTGGTAT TCTTCAGCCT
 481 GCGTGTTGTT AACATGCCGT TCTCCTATGA CCTGTTCAAC AAGAGTTCTC TGGAGTATCA
 541 AGCCCTGGAA CAACGATTCA CAGACCTGCT GGTTCCCTAT CWACGATCGA ATCTTACGGG
 601 ATTTAAGCAA CTGGAAATAC TCAGCTTCAG AAACGGAAGT GTGATCGTGA ACAGCAAAGT
 661 GCGGTTTGCA AAGGCGGTAC CCTACAACCT CACCCAGGCC GTGCGCGGGG TCTTGGAGGA
 721 TCTTCGGTCC ACCGCAGCTC AAGGGCTCAA TCTGGAAATC GAAAGCTACT CCCTCGACAT
 781 TGAACCAGCT GATCAGGCGG ATCCCTGCAA ACTTCTAGAC TGTGGCAAAT TTGCCCAGTG
 841 TGTAAAGAAT GAGTGGACAG AGGAAGCAGA GTGTCGCTGC AGACAGGGAC ATGAGAGCCA
 901 CGGGACCCTG GACTACCAGA CCCTGAACCT CTGTCCCCCT GGAAAGACTT GTGTGGCCGG
 961 CCGAGAACAA GCAACTCCAT GCAGGCCACC AGATCACTCT ACAAACCAAG CTCAGGAACC
1021 TGGTGTTAAA AAGCTACGTC AGCAAAATAA GGTAGTCAAG AAACGAAATT CTAAACTATC
1081 AGCTATAGGA TTTGAAAAAT TTGAAGACCA GGACTGGGAG GGAAATTAAA AGCTGGAATC
1141 ATATGCATTA TGTTGCAAAC TCTGTTGAAA GGAAACTTTA TTTCTTAAAG AAAGGTGTAT
1201 CTGTTCTGTT AACTTCTGAA AAACAGAGGG AGAGATTCAG TGGTCATTGG AATACAGGCA
1261 TGTAATCAAC TTTGAGACTC AGCATGCTTG AACAAGAGCA CAGGCGTGTA TTTGATAAGC
1321 C
//
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SEQ ID NO:13 Mouse IPM 150 amino acid sequence, isoform E

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1 MNFQIKHAIF VFGIFLQVQG IKDTSIKIFS SEIKNIDKTP RIETIESTST VHKVSTMKRQ 61 PCQKYQN //
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SEQ ID NO:14 Monkey IPM 150 cDNA (partial)

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1 ATTTTCTTTC CGAACGGGGT TAAAGTCTGT CCACAGGAAT CCATGAAACA GATTTTAGCC 61 AGTCTTCAAG CTTATTATAG ATTGAGAGTG TGTCAGGAAG CAGTATGGGA AGCATATCGG 121 ATCTTTCTGG ATCGCATCCC TGACACAGGG GAATATCAGG ACTGGGTCAG CTTCTGCCAG 181 CAGGAGACCT TCTGCCTCTT TGACATCGGA CAAAACTTCA GCAATTCCCA GGAGCACCTG 241 GATCTTCTCC AGCAGAGAAT AAAACAGAGA AGTTTCCCTG AGAGAAAAGA TGAAGTATCT 301 ACAGAGAAGA CATTGGGAGA GCCTAGTGAA ACCATTGTGG TTTCAACAGA TGTTGCCAGC 361 GTCTCACTTG GGCCTTTCCC TGTCACTCCT GATGACACCC TCCTCAATGA AATTCTCGAT 421 AATGCACTCA ACGACACCAA GATGCCTACA ACAGAAAGAG AAACAGAACT CGCTGTGTCT 481 GAGGAGCAGA GGGTGGAGCT CAGCATCTCT CTGATAAACC AGAGGTTCAA GGCAGAGCTC 541 GCTGACTCTC AGTCA
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SEQ ID NO:15 Monkey IPM 150 amino acid sequence (partial)

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1 IFFPNGVKVC PQESMKQILA SLQAYYRLRV CQEAVWEAYR IFLDRIPDTG EYQDWVSFCQ 61 QETFCLFDIG QNFSNSQEHL DLLQQRIKQR SFPERKDEVS TEKTLGEPSE TIVVSTDVAS 121 VSLGPFPVTP DDTLLNEILD NALNDTKMPT TERETELAVS EEQRVELSIS LINQRFKAEL 181 ADSQS
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SEQ ID NO:16 Human IPM 200 (isoform A) cDNA

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1 CGGGCTACTT TGAAAGGACA ACCATTTTTC TTTCCGCTAA TTTATAATGG TTTTGAAGTG
  61 GTTGTTCATT CTCAAACATA GACTTTTAAA TGTTAGGTCT TTCCTATAAC TCTTTGTTAT
 181 TTTTGGCCCA AATGATTATG TTTCCTCTTT TTGGGAAGAT TTCTCTGGGT ATTTTGATAT 241 TTGTCCTGAT AGAAGGAGAC TTTCCATCAT TAACAGCACA AACCTACTTA TCTATAGAGG
 301 AGATCCAAGA ACCCAAGAGT GCAGTTTCTT TTCTCCTGCC TGAAGAATCA ACAGACCTTT
 361 CTCTAGCTAC CAAAAAGAAA CAGCCTCTGG ACCGCAGAGA AACTGAAAGA CAGTGGTTAA
 421 TCAGAAGGCG GAGATCTATT CTGTTTCCTA ATGGAGTGAA AATCTGCCCA GATGAAAGTG
 481 TTGCAGAGGC TGTGGCAAAT CATGTGAAGT ATTTTAAAGT CCGAGTGTGT CAGGAAGCTG 541 TCTGGGAAGC CTTCAGGACT TTTTGGGATC GACTTCCTGG GCGTGAGGAA TATCATTACT
 601 GGATGAATTT GTGTGAGGAT GGAGTCACAA GTATATTTGA AATGGGCACA AATTTTAGTG
 661 AATCTGTGGA ACATAGAAGC TTAATCATGA AGAAACTGAC TTATGCAAAG GAAACTGTAA
721 GCAGCTCTGA ACTGTCTTCT CCAGTTCCTG TTGGTGATAC TTCAACATTG GGAGACACTA
 781 CTCTCAGTGT TCCACATCCA GAGGTGGACG CCTATGAAGG TGCCTCAGAG AGCAGCTTGG
 841 AAAGGCCAGA GGAGAGTATT AGCAATGAAA TTGAGAATGT GATAGAAGAA GCCACAAAAC
 901 CAGCAGGTGA ACAGATTGCA GAATTCAGTA TCCACCTTTT GGGGAAGCAG TACAGGGAAG
961 AACTACAGGA TTCCTCCAGC TTTCACCACC AGCACCTTGA AGAAGAATTT ATTTCAGAGG
1021 TTGAAAATGC ATTTACTGGG TTACCAGGCT ACAAGGAAAT TCGTGTACTT GAATTTAGGT
1081 CCCCCAAGGA AAATGACAGT GGCGTAGATG TTTACTATGC AGTTACCTTC AATGGTGAGG
1141 CCATCAGCAA TACCACCTGG GACCTCATTA GCCTTCACTC CAACAAGGTG GAAAACCATG
1201 GCCTTGTGGA ACTGGATGAT AAACCCACTG TTGTTTATAC AATCAGTAAC TTCAGAGATT
1261 ATATTGCTGA GACATTGCAG CAGAATTTTT TGCTGGGGAA CTCTTCCTTG AATCCAGATC
1321 CTGATTCCCT GCAGCTTATC AATGTGAGAG GAGTTTTGCG TCACCAAACT GAAGATCTAG
1381 TTTGGAACAC CCAAAGTTCA AGTCTTCAGG CAACGCCGTC ATCTATTCTG GATAATACCT
1441 TTCAAGCTGC ATGGCCCTCA GCAGATGAAT CCATCACCAG CAGTATTCCA CCACTTGATT
1501 TCAGCTCTGG TCCTCCCTCA GCCACTGGCA GGGAACTCTG GTCAGAAAGT CCTTTGGGTG
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1561 ATTTAGTGTC TACACACAAA TTAGCCTTTC CCTCGAAGAT GGGCCTCAGC TCTTCCCCAG 1621 AGGTTTTAGA GGTTAGCAGC TTGACTCTTC ATTCTGTCAC CCCGGCAGTG CTTCAGACTG 1681 GCTTGCCTGT GGCTTCTGAG GAAAGGACTT CTGGATCTCA CTTGGTAGAA GATGGATTAG 1741 CCAATGTTGA AGAGTCAGAA GATTTTCTTT CTATTGATTC ATTGCCTTCA AGTTCATTCA 1801 CTCAACCTGT GCCAAAAGAA ACAATACCAT CCATGGAAGA CTCTGATGTG TCCTTAACAT 1861 CTTCACCATA TCTGACCTCT TCTATACCTT TTGGCTTGGA CTCCTTGACC TCCAAAGTCA 1921 AAGACCAATT AAAAGTGAGC CCTTTCCTGC CAGATGCATC CATGGAAAAA GAGTTAATAT 1981 TTGACGGTGG TTTAGGTTCA GGGTCTGGGC AAAAGGTAGA TCTGATTACT TGGCCATGGA 2041 GTGAGACTTC ATCAGAGAAG AGCGCCGAAC CACTGTCCAA GCCGTGGCTT GAAGATGATG 2101 ATTCACTTTT GCCAGCTGAG ATTGAAGACA AGAAACTAGT TTTAGTTGAC AAAATGGATT 2161 CCACAGACCA AATTAGTAAG CACTCAAAAT ATGAACATGA TGACAGATCC ACACACTTTC 2221 CAGAGGAAGA GCCTCTTAGT GGGCCTGCTG TGCCCATCTT CGCAGATACT GCAGCTGAAT 2281 CTGCGTCTCT AACCCTCCCC AAGCACATAT CAGAAGTACC TGGTGTTGAT GATTGCTCAG 2341 TTACCAAAGC ACCTCTTATA CTGACATCTG TAGCAATCTC TGCCTCTACT GATAAATCAG 2401 ATCAGGCAGA TGCCATCCTA AGGGAGGATA TGGAACAAAT TACTGAGTCA TCCAACTATG 2461 AATGGTTTGA CAGTGAGGTT TCAATGGTAA AGCCAGATAT GCAAACTTTG TGGACTATAT 2521 TGCCAGAATC AGAGAGAGTT TGGACAAGAA CTTCTTCCCT AGAGAAATTG TCCAGAGACA 2581 TATTGGCAAG TACACCACAG AGTGCTGACA GGCTCTGGTT ATCTGTGACA CAGTCTACCA 2641 AATTGCCTCC AACCACAATC TCCACCCTGC TAGAGGATGA AGTAATTATG GGTGTACAGG 2701 ATATTTCGTT AGAACTGGAC CGGATAGGCA CAGATTACTA TCAGCCTGAG CAAGTCCAAG 2761 AGCAAAATGG CAAGGTTGGT AGTTATGTGG AAATGTCAAC AAGTGTTCAC TCCACAGAGA 2821 TGGTTAGTGT GGCTTGGCCC ACAGAAGGAG GAGATGACTT GAGTTATACC CAGACTTCAG 2881 GAGCTTTGGT GGTTTTCTTC AGCCTCCGAG TGACTAACAT GATGTTTTCA GAAGATCTGT 2941 TTAATAAAAA CTCCTTGGAG TATAAAGCCC TGGAGCAAAG ATTCTTAGAA TTGCTGGTTC 3001 CCTATCTCCA GTCAAATCTC ACGGGGTTCC AGAACTTAGA AATCCTCAAC TTCAGAAATG 3061 GCAGCATTGT GGTGAACAGT CGAATGAAGT TTGCCAATTC TGTCCCTCCT AACGTCAACA 3121 ATGCGGTGTA CATGATTCTG GAAGACTTTT GTACCACTGC CTACAATACC ATGAACTTGG 3181 CTATTGATAA ATACTCTCTT GATGTGGAAT CAGGTGATGA AGCCAACCCT TGCAAGTTTC 3241 AGGCCTGTAA TGAATTTTCA GAGTGTCTGG TCAACCCCTG GAGTGGAGAA GCAAAGTGCA 3301 GATGCTTCCC TGGATACCTG AGTGTGGAAG AACGGCCCTG TCAGAGTCTC TGTGACCTAC 3361 AGCCTGACTT CTGCTTGAAT GATGGAAAGT GTGACATTAT GCCTGGGCAC GGGGCCATTT 3421 GTAGGTGCCG GGTGGGTGAG AACTGGTGGT ACCGAGGCAA GCACTGTGAG GAATTTGTGT 3481 CTGAGCCCGT GATCATAGGC ATCACTATTG CCTCCGTGGT TGGACTTCTT GTCATCTTTT 3541 CTGCTATCAT CTACTTCTTC ATCAGGACTC TTCAAGCACA CCATGACAGG AGTGAAAGAG 3601 AGAGTCCCTT CAGTGGCTCC AGCAGGCAGC CTGACAGCCT CTCATCTATT GAGAATGCTG 3661 TGAAGTACAA CCCCGTGTAT GAAAGTCACA GGGCTGGATG TGAGAAGTAT GAGGGACCCT 3721 ATCCTCAGCA TCCCTTCTAC AGCTCTGCTA GCGGAGACGT GATTGGTGGG CTGAGCAGAG 3781 AAGAAATCAG ACAGATGTAT GAGAGCAGTG AGCTTTCCAG AGAGGAAATT CAAGAGAGAA 3841 TGAGAGTTTT GGAACTGTAT GCCAATGATC CTGAGTTTGC AGCTTTTGTG AGAGAGCAAC 3901 AAGTGGAAGA GGTTTAACCA AAACTCCTGT TCTGAAACTG ATTAGAAGCC TGGAGAAGAT 3961 GGAGATTACT TGTTACTTAT GTCATATAAT TAACCTGGAT TTTAAACACT GTTGGAAGAA 4021 GAGTTTTCTA TGAAAAAATT AAATATAGGG CACACTGTTT TTTTTTCAGC TTAAGTTTTC 4081 AGAATGTAGT AAGAGATGTT ACCATTTTTA TTTCTATAAA GACTGAATGC TGTGTTTAAA 4141 TAAATTGAAA ACTACGTAAA AAAAAA 11

SEQ ID NO:17 Human IPM 200 amino acid sequence, isoform A

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1 MIMFPLFGKI SLGILIFVLI EGDFPSLTAQ TYLSIEEIQE PKSAVSFLLP EESTDLSLAT
61 KKKQPLDRRE TERQWLIRRR RSILFPNGVK ICPDESVAEA VANHVKYFKV RVCQEAVWEA
121 FRTFWDRLPG REEYHYWMNL CEDGVTSIFE MGTNFSESVE HRSLIMKKLT YAKETVSSSE
181 LSSPVPVGDT STLGDTTLSV PHPEVDAYEG ASESSLERPE ESISNEIENV IEEATKPAGE
241 QIAEFSIHLL GKQYREELQD SSSFHHQHLE EEFISEVENA FTGLPGYKEI RVLEFRSPKE
301 NDSGVDVYYA VTFNGEAISN TTWDLISLHS NKVENHGLVE LDDKPTVVYT ISNFRDYIAE
361 TLQQNFLLGN SSLNPDPDSL QLINVRGVLR HQTEDLVWNT QSSSLQATPS SILDNTFQAA
421 WPSADESITS SIPPLDFSSG PPSATGRELW SESPLGDLVS THKLAFPSKM GLSSSPEVLE
481 VSSLTLHSVT PAVLQTGLPV ASEERTSGSH LVEDGLANVE ESEDFLSIDS LPSSSFTQPV
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541 PKETIPSMED SDVSLTSSPY LTSSIPFGLD SLTSKVKDQL KVSPFLPDAS MEKELIFDGG
601 LGSGSGQKVD LITWPWSETS SEKSAEPLSK PWLEDDDSLL PAEIEDKKLV LVDKMDSTDQ
661 ISKHSKYEHD DRSTHFPEEE PLSGPAVPIF ADTAAESASL TLPKHISEVP GVDDCSVTKA
721 PLILTSVAIS ASTDKSDQAD AILREDMEQI TESSNYEWFD SEVSMVKPDM QTLWTILPES
781 ERVWTRTSSL EKLSRDILAS TPQSADRLWL SVTQSTKLPP TTISTLLEDE VIMGVQDISL
841 ELDRIGTDYY QPEQVQEQNG KVGSYVEMST SVHSTEMVSV AWPTEGGDDL SYTQTSGALV
901 VFFSLRVTNM MFSEDLFNKN SLEYKALEQR FLELLVPYLQ SNLTGFQNLE ILNFRNGSIV
961 VNSRMKFANS VPPNVNNAVY MILEDFCTTA YNTMNLAIDK YSLDVESGDE ANPCKFQACN
1021 EFSECLVNPW SGEAKCRCFP GYLSVEERPC QSLCDLQPDF CLNDGKCDIM PGHGAICRCR
1081 VGENWWYRGK HCEEFVSEPV IIGITIASVV GLLVIFSAII YFFIRTLQAH HDRSERESPF
1141 SGSSRQPDSL SSIENAVKYN PVYESHRAGC EKYEGPYPQH PFYSSASGDV IGGLSREEIR
1201 QMYESSELSR EEIQERMRVL ELYANDPEFA AFVREQQVEE V

SEQ ID NO:18 Human IPM 200 cDNA sequence, isoform C

1 AAACTTAAGC TTGGAGTTTG GAAGTTTCAA GGATTTGGAC ACTCAATTAA GGATTCTGTC 61 CTCTCCTCAT TCCTTTGGTT TTGGCCCAAA TGATTATGTT TCCTCTTTTT GGGAAGATTT 121 CTCTGGGTAT TTTGATATTT GTCCTGATAG AAGGAGACTT TCCATCATTA ACAGCACAAA 181 CCTACTTATC TATAGAGGAG ATCCAAGAAC CCAAGAGTGC AGTTTCTTTT CTCCTGCCTG 241 AAGAATCAAC AGACCTTTCT CTAGCTACCA AAAAGAAACA GCCTCTGGAC CGCAGAGAAA 301 CTGAAAGACA GTGGTTAATC AGAAGGCGGA GATCTATTCT GTTTCCTAAT GGAGTGAAAA 361 TCTGCCCAGA TGAAAGTGTT GCAGAGGCTG TGGCAAATCA TGTGAAGTAT TTTAAAGTCC 421 GAGTGTGTCA GGAAGCTGTC TGGGAAGCCT TCAGGACTTT TTGGGATCSA CTTCCTGGGC 481 GTGAGGAATA TCATTACTGG ATGAATTTGT GTGAGGATGG AGTCACAAGT ATATTTGAAA 541 TGGGCACAAA TTTTAGTGAA TCTGTGGAAC ATAGAAGCTT AATCATGAAG AAACTGACTT 601 ATGCAAAGGA AACTGTAAGC AGCTCTGAAC TGTCTTCTCC AGTTCCTGTT GGTGATACTT 661 CAACATTGGG AGACACTACT CTCAGTGTTC CACATCCAGA GGTGGACGCC TATGAAGGTG 721 CCTCAGAGAG CAGCTTGGAA AGGCCAGAGG AGAGTATTAG CAATGAAATT GAGAATGTGA 781 TAGAAGAAGC CACAAAACCA GCAGGTGAAC AGATTGCAGA ATTCAGTATC CACCTTTTGG 841 GGAAGCAGTA CAGGGAAGAA CTACAGGATT CCTCCAGCTT TCACCACCAG CACCTTGAAG 901 AAGAATTTAT TTCAGAGGTT GAAAATGCAT TTACTGGGTT ACCAGGCTAC AAGGAAATTC 961 GTGTACTTGA ATTTAGGTCC CCCAAGGAAA ATGACAGTGG CGTAGATGTT TACTATGCAG 1021 TTACCTTCAA TGGTGAGGCC ATCAGCAATA CCACCTGGGA CCTCATTAGC CTTCACTCCA 1081 ACAAGGTGGA AAACCATGGC CTTGTGGAAC TGGATGATAA ACCCACTGTT GTTTATACAA 1141 TCAGTAACTT CAGAGATTAT ATTGCTGAGA CATTGCAGCA GAATTTTTTG CTGGGGAACT 1201 CTTCCTTGAA TCCAGATCCT GATTCCCTGC AGCTTATCAA TGTGAGAGGA GTTTTGCGTC 1261 ACCAAACTGA AGATCTAGTT TGGAACACCC AAAGTTCAAG TCTTCAGGCA ACGCCGTCAT 1321 CTATTCTGTG CTTCARACTG GCTTGCCTGT GGCTTCTGAG GAAAGGACTT CTGGATCTCA 1381 CTTGGTAGAA GATGGATTAG CCAATGTTGA AGAGTCAGAA GATTTTCTTT CTATTGATTC 1441 ATTGCCTTCA AGTTCATTCA CTCAACCTGT GCCAAAAGAA ACAATACCAT CCATGGAAGA 1501 CTCTGATGTG TCCTTAACAT CTTCACCATA TCTGACCTCT TCTATACCTT TTGGCTTGGA 1561 CTCCTTGACC TCCAAAGTCA AAGACCAATT AAAAGTGAGC CCTTTCCTGC CAGATGCATC 1621 CATGGAAAAA GAGTTAATAT TTGACGGTGG TTTAGGTTCA GGGTCTGGGC AAAAGGTAGA 1681 TCTGATTACT TGGCCATGGA GTGAGACTTC ATCAGAGAAG AGCGCTGAAC CACTGTCCAA 1741 GCCGTGGCTT GAAGATGATG ATTCACTTTT GCCAGCTGAG ATTGAAGACA AGAAACTAGT 1801 TTTAGTTGAC AAAATGGATT CCACAGACCA AATTAGTAAG CACTCAAAAT ATGAACATGA 1861 TGACAGATCC ATACACTTTC CAGAGGAAGA GCCTCTTAGT GGGCCTGCTG TGCCCATCTT 1921 CGCAGATACT GCAGCTGAAT CTGCGTCTCT AACCCTCCCC AAGCACATAT CAGAAGTACC 1981 TGGTGTTGAT GATTACTCAG TTACCAAAGC ACCTCTTATA CTGACATCTG TAGCAATCTC 2041 TGCCTCTACT GATAAATCAG ATCAGGCAGA TGCCATCCTA AGGGAGGATA TGGAACAAAT 2101 TACTGAGTCA TCCAACTATG AATGGTTTGA CAGTGAGGTT TCAATGGTAA AGCCAGATAT 2161 GCAAACTTTG TGGACTATAT TGCCAGAATC AGAGAGAGTT TGGACAAGAA CTTCTTCCCT 2221 AGAGAAATTG TCCAGAGACA TATTGGCAAG TACACCACAG AGTGCTGACA GGCTCTGGTT 2281 ATCTGTGACA CAGTCTACCA AATTGCCTCC AACCACAATC TCCACCCTGC TAGAGGATGA 2341 AGTAATTATG GGTGTACAGG ATATTTCGTT AGAACTGGAC CGGATAGGCA CAGATTACTA

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2401 TCAGCCTGAG CAAGTCCAAG AGCAAAATGG CAAGGTTGGT AGTTATGTGG AAATGTCAAC
2461 AAGTGTTCAC TCCACAGAGA TGGTTAGTGT GGCTTGGCCC ACAGAAGGAG GAGATGACTT
2521 GAGTTATACC CAGACTTCAG GAGCTTTGGT GGTTTTCTTC AGCCTCCGAG TGACTAACAT
2581 GATGTTTTCA GAAGATCTGT TTAATAAAAA CTCCTTGGAG TATAAAGCCC TGGAGCAAAG
2641 ATTCTTAGAA TTGCTGGTTC CCTATCTCCA GTCAAATCTC ACGGGGTTCC AGAACTTAGA
2701 AATCCTCAAC TTCAGAAATG GCAGCATTGT GGTGAACAGT CGAATGAAGT TTGCCAATTC
2761 TGTCCCTCCT AACGTCAACA ATGCGGTGTA CATGATTCTG GAAGACTTTT GTACCACTGC
2821 CTACAATACC ATGAACTTGG CTATTGATAA ATACTCTCTT GATGTGAAT CAGGTGATGA
2881 AGCCAACCCT TGCAAGTTTC AGGCCTGTAA TGAATTTTCA GAAGTGTCTG GTCAACCCCT
2941 GGAGTGGAGA AACAAAAGTG CAGA
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SEQ ID NO:19 Human IPM 200 amino acid sequence, isoform C

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1 MIMFPLFGKI SLGILIFVLI EGDFPSLTAQ TYLSIEEIQE PKSAVSFLLP EESTDLSLAT
61 KKKQPLDRRE TERQWLIRRR RSILFPNGVK ICPDESVAEA VANHVKYFKV RVCQEAVWEA
121 FRTFWDXLPG REEYHYWMNL CEDGVTSIFE MGTNFSESVE HRSLIMKKLT YAKETVSSSE
181 LSSPVPVGDT STLGDTTLSV PHPEVDAYEG ASESSLERPE ESISNEIENV IEEATKPAGE
241 QIAEFSIHLL GKQYREELQD SSSFHHQHLE EEFISEVENA FTGLPGYKEI RVLEFRSPKE
301 NDSGVDVYYA VTFNGEAISN TTWDLISLHS NKVENHGLVE LDDKPTVVYT ISNFRDYIAE
361 TLQQNFLLGN SSLNPDPDSL QLINVRGVLR HQTEDLVWNT QSSSLQATPS SILCFXLACL
421 WLLRKGLLDL TW
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SEQ ID NO:20 Human IPM 200 cDNA sequence, isoform F

SEQ ID NO:21 Human IPM 200 amino acid sequence, isoform F

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1 MIMFPLFGKI SLGILIFVLI EGDFPSLTAQ TYLSIEEIQA L
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SEQ ID NO:22 Human IPM 200, regulatory region

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1 GAACACTTGT AATACAAAAC AATTCCTATT TACAAAGTTT ACTGGTAATA CAAATACAGT
61 AGTTTACAGA GAACTTTCAT GTCTCTTAAT TCTTAACAAC GACCCTGTGA TACAGGTAGA
121 GATTATCACA TGTAATTTCT TTGGTGAGTA AACCGGCTCA AAGAGCTTAG GTTATTTACC
181 AAAATCAAAT ATTAAGTGAT AAAACCAAGA TTTGAGTCCA GGGTTTCTCA ATCTTAAATA
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241 CAGGAATCTT TCTAGATTAC TATGATTCTC AGAAGTTTTT TTTAGCTTTT TGGTCAAGGC 301 TGTCAAAAAG AATAATTGCC AACTTAATAT TTGTTACCTA AGAGTTGTCC CTTGTTCTGA 361 ATTGTCAATA TGAAGCTTTT CTTAAGATTA AACTTTGACT CAGCTAATAA AATTTTCGGC 421 TTTTTTCTCC TACTCATACA ATAAATTTGG CAAGTAAGTT TCTTATAAGC TTACCAGTAT 481 TTTGCAAATA CAACTATGCA AATATATTTA ATGGTCATTT AGGTTTATTA GCTTTTATAA 541 AGGCTGAAAA TGTGGTTTAT TTGAGGCTGT ATTGAAAAAA TATACTTGAG CTTTTCCTAA 601 AGCATAAAAT AACATTGAGG GTGATTTAGC TAACACAATT AGTCAAGGAT TCTCAAGAGG 661 AATGTGGTTT AGATCTTTAC AATACACTTT TTTTCAGAGA ATTTTGCCAG AGATAACATG 721 AAATAAAATA TAATTTCATT GCTATTTGAT AGTAAATCCA AGCTTCCACA GGGATTCTGA 781 TGAATTGCTT TCTACTAGGT TTACTTGATT TAAAAAACTG TTCTAATATA GAGAATTTCA 841 TCTGCAGGGA AAATGTTTTC TTGGTTAAGA GTTCCTCATG TAGATAAACA CACTGGGCCT 901 CACATTTAAT GGCAAATTAA GCAACAAAGT TATCGCACAG CTATCATTTA TATTAAGTGC 961 TTAATATGTT CCGGGCACTA CTCTAAGCAA AGTGAAGATT GAATTAGTTA ATTAGTTAAT 1021 TTAATCCTCA CATTAGCTCT ACCATGAGTT TACTATTTCT ATTCCATTTT ATACGTAAGG 1081 AAGGAGACAA AGTAAGTGAT TTTTCTATCA AGGAAGGAAA TTTGCAAGAG AATAGTTTCA 1141 TTACAAAAC TAAATTTGTA CGTAGCTCTG TATTATTGAA ATAGGTAGAT ATAGTCAGTC 1201 TGGACTTTTT ATGCTTATAC ATCTTAGTCC CTAGGAAAAC CCAGAACTAA CAGATTCAGA 1261 AAAGTTGGAA AAATCAGTGA ATTATATGTG AAACACATTA TTCTTAGTGG ACTGCTTGTT 1321 AAAGGCAAGG AGAGTGTTAG TAAAGAGCTT AGGTAGATTA GAATAAAGAA ATTGTCTCTC 1381 TCCATCTGCT CTAATTAGCT TATCTCACCA GCTTTTATAG CATGCTGGTT ATTTCAGAAA 1441 AGAAGTGAGA GCTACTTTGA AAGGACAACC ATTTTTCTTT CCGCTAATTT ATAATGGTTT 1501 TGAAGTGGTT GTTCATTCTC AAACATAGAC TTTTAAATGT TAGGTCTTTC CTATAACTCT 1561 TTGTTATTGG AAGTTTCAAG GATTTGGACA CTCAATTAAG GATTCTGTCC TCTCCTCATT 1621 CCTTTGGTTT TGGCCCAAAT GATTATGTTT CCTCTTTTTG GGAAGATTTC TCTGGGTATT 1681 TTGATATTTG TCCTGATAGA AGGAGACTTT CCATCATTAA CAGGTATTTA AAAATCTACA 1741 TTTGTTTGTA TCTTTCCATA TCTGTAGTAT ATGTTCTTCA AAAATAGGAT TATTTGATGT 1801 GATTGCTGTA AGAAATGGAA TCAAATACTT TATTAATCTT TGATATGGCT TCATTTAAAC 1861 CGTTTTAAAA TATCTCCCAA TAATTTTGGT TTTCCCTCAT TAGTAATTTC TGGTTTAAAC 1921 CTTACTTTTA TTTATTTTGT TGAAATTGGA TGTGTATTTA CTTGATTTTG ATAACAATCT 1981 TGAATGAAAG GAGTGGGAGT TAAATGGAAA AAGATGGACT GCCTCACTCC TCTTTTCCTT 2041 AGATATGCAT GCCTGCCTAT GATTTGGGCA CTGGCTTCTC TATCTTAATG TAGCCCAAGT 2101 GTCAGTTTTT CTTTAGTTGT TACCTTTTGT ACTGTATCTT CATTATCGAA GACTTGACTA 2161 TACTTTCACT CTGTAGCACA AACCTACTTA TCTATAGAGG AGATCCAAGA ACCCAAGAGT 2221 GCAGTTTCTT TTCTCCTGCC TGAAGAATCA ACAGACCTTT CTCTAGCTAC CAAAAAGAAA 2281 CAGCCTCTGG ACCGCAGAGA AACTGAAAGA CAGTGGTTAA TCAGAAGGCG GAGATCTATT 2341 CTGTTTCCTA ATGGAGTGAA AATCTGCCCA GATGAAAGTG TTGCAGAGGC TGTGGCAAAT 2401 CATGTGAAGT ATTTTAAAGT CCGAGGTAAG CGAACATCCA AATCCTTCAG CTCCATAATG 2461 AAATTCAAAC ATAGTTTAAT CATTTGTTAG GTAACATTGT AAATCAAAAT TTATGATAAT 2521 TTAGACAGGA CTGAGCCAAA ACTACCTTTC TACTGTTAAG AATATAGTGT TAATGGTAAC 2581 TTCAGAGAAC AGTTTACATT AAGAGAGGAG GTTTGTTTTT TTTCCAGTGC CCTCCAGTTA 2641 AGGCAATAAT ATCATTTAAT AATGACATGC ACTTTGAACC AAAGGAAGAA CGCTTTCATG 2701 ATTTGAGTTT GTAGCTTTTG GTGCGTTATG TAAGAAACTT TTTTCACATG AGGGCAGTCA 2761 CAATAAGATG TCTTTCATTA ATTTCAACAA CATATTCAGA GAGGAAATGT CTTAAATCTT 2821 TTTAAGCACT TCAAAAATAC CAGTTTATGT TTTGGGCTAC ATTAATTTTA ATTTTTACTT 2881 CTTCATTACA GTAAATGCCT AAGTWTACCG ACAAAATAGC TTTACCAAAG NTATACTCAC 2941 CTGCTTGCCT ATTTAATTAA TAGTTATTAT ATATACAAAT ATAATGTTTC TATATTTTAT 3001 AGTTTAGATA T

SEQ ID NO:23 Mouse IPM 200 cDNA sequence (partial)

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1 CCGTGGCAAA CCATGTGAAG TATTTTAAAG CCCGAGTGTG CCAGGAAGCC ATCTGGGAAG
61 CCTTCAGGAC GTTTTGGGAT CGACTTCCTG GGCGTGATGA ATATCGTCAC TGGATGAATT
121 TATGTGAGGA TGGAGTCACA AGTGTATTTG AAATGGGCGC CCATTTTAGT CAGTCTGTGG
181 AACATAGAAA CCTAATCATG AAGAAACTGG CTTACACAAG GGAAGCTGAG AGCAGCTCCT
241 GCAAGGATCA GTCCTGTGGG CCTGAGTTGT CCTTTCCAGT TCCTATTGGT GAGACCTCAA
301 CACTGACAGG TGCTGTCTCC AGTGCTTCCT ATCCAGGGTT GGCTTCGGAG AGCAGCGCAG
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361 CGTCACCGCA GGAGAGTATC AGCAATGAAA TTGAGAATGT GACAGAGGAG CCCACACAAC 421 CAGCTGCTGA ACAGATTGCG GAATTCAGCA TCCAACTTCT GGGGAAGCGA TACAGTGAAG 481 AACTGCGGGA TCCCTCCAGC GCCCTCTACC GGCTCCTCGT GGAAGAGTTT ATTTCAGAGG 541 TTGAAAAAGC ATTCACAGGG TTACCTGGCT ACAAGGGCAT CCGTGTTCTG GAATTCAGGG 601 CCCCGGAGGA AAATGACAGT GGGATAGATG TTCACTATGC AGTTACCTTC AATGGCGAAG 661 CCATCAGCAA TACCACCTGG GACCTCATAA GCCTTCACTC CAACAAGGTA GAAAACCATG 721 GCCTTGTAGA GATGGATGAT AAACCCACTG CTGTCTATAC AATTAGTAAC TTCAGAGATT 781 ATATCGCTGA GACGCTGCAC CAGAACTTTT TGATGGGAAA TTCCTCTTTG AATCCAGATC 841 CCAAGCCTCT CCAGCTCATC AATGTGAGAG GAGTTTTGCT CCCCCAAACA GAAGACATAG 901 TTTGGAACAC CCAAAGTTCA AGTCTTCAGG TGACAACATC CTCTATTTTN GTGCTTCAGC 961 CTGACCTGCC TGTGGCTCCT GAGGGAAGGA CTTCTGGATC GTTCATATTA GAAGATGGGT 1021 TAGCCAGCAC TGAAGAATTA GAAGATACTT CTATTGATGG ATTGCCTTCA AGCCCATTAA 1081 TTCAACCTGT GCCAAAAGAA ACAGTACCAC CTATGGAAGA CTCTGACACG GCTCTCTTGT 1141 CCACACCACA TCTGACCTCT TCTGCTATAG AAGACCTTAC TAAAGACATA GGGACACCTT 1201 CTGGCTTGGA GTCCTTGGCT TCAAACATCT CAGACCAGTT GGAAGTGATC CCATGGTTTC 1261 CAGACACCTC TGTGGAAAAA GACTTCATTT TTGAAAGTGG CTTGGGTTCT GGGTCTGGGA 1321 AAGATGTAGA TGTGATTGAT TGGCCATGGA GTGAGACTTC ATTAGAGAAG ACCACTAAAC 1381 CACTGTCAAA GTCATGGTCT GAAGAACAGG ATGCACTATT ACCAACTGAG GGTAGAGAAA 1441 AATTACATAT AGATGGCAGA GTAGATTCCA CAGAACAAAT TATTGAATCA TCAGAACATA 1501 GATATGGAGA TAGGCCCATA CATTTTATAG AGGAAGANTC CCATGTTAGA TCTACTATAC 1561 CCATCTTTGT AGAGTCCGCA ACTCCACCTA CATCTCCAAT CTTTTCAAAA CACACTTCAG 1621 ATGTACCAGA CATTGATTCT TACTCACTTA CCAAACCACC CTTCTTACCG GTAACTATAG 1681 CAATCCCTGC TTCCACTAAG AAAACAGATG AGGTACTCAA GGAAGATATG GTACATACAG 1741 AATCATCCAG TCACAAAGAA CTTGACAGTG AGGTTCCAGT GTCAAGGCCA GATATGCAGC 1801 CTGTGTGGAC CATGTTGCCA GAATCAGATA CAGTTTGGAC AAGAACTTCT TCCTTAGGGA 1861 AATTGTCCAG AGACACATTG GCAAGTACAC CAGAGAGCAC TGACAGACTC TGGTTGAAAG 1921 CTTCCATGAC ACAGTCCACT GAATTGCCTT CAACCACCCA CTCCACCCAG CTAGAGGAGG 1981 AAGTAATAAT GGCGGTCCAG GATATTTCAT TAGAACTAGA TCAGGTAGGC ACAGATTATT 2041 ATCAGTCCGA GCTAACTGAA GAACAACATG GCAAGGCTGA CAGCTATGTG GAAATGTCTA 2101 CCAGTGTTCA CTACACAGAG ATGCCTATTG TGGCTCTGCC CACAAAAGGA GGTGTTCTGA 2161 GTCACACCCAG ACTGCAGGAG CATTGGTGGT TTTCTTCAGC CTCCGCGTGA CAAACATGT 2221 TGTTTTCAGAA GACTTGTTTA ACAAAAACTC TTTGGAATAT AAAGCCCTGG AACAAAGAT 2281 TCTTAGAACTG CTGGCTCCCT ATCTCCAGTC AAATCTGTCA GGGTTCCAGA ACCTAGAAA 2341 TCCTGAGTTTC AGAAACGGCA GCATTGTGGT GAACAGCCGA GTGAGGTTCG CCGAGTCTG 2401 CCCCTCCTAAT GTCAACAAGG CCATGTATAG GATTCTGGAA GACTTTTGTA CCACTGCCT 2461 ACCAAACCATG AACTTGGATA TCGATAAGTA CTCCCTGGAC GTGGAATCAG GTGATGAGG 2521 CCAACCCTTGC AAGTTTCAGG CCTGTAATGA ATTTTCTGAG TGTTTGGTAA ATCCATGGA 2581 GTGGAGAAGCA AAGTGCAAAT GCTACCCTGG GTACCTGAGT GTGGATGAAC TGCCTTGTC 2641 AAAGTCTCTGT GATCTACAGC CTGACTTCTG CTTGAACGAT GGAAAGTGTG ACATTATGC 2701 CTGGGCATGGA GCCATTTGTA GATGCCGGGT TGGTTCAAAC TGGTGGTATC GAGGCCAAC 2761 ACTGTGAGGAG TTTGTGTCTG AGCCCTTTGT CATAGGCATC ACTATAGCCT CTGTGGTTA 2821 GCTTTCTCCTT GTTGCTTCTG CTGTCGTCTT CTTCCTTGTG AAGATGCTTC AAGCTCAGA 2881 ATGTCAGGAGA GAAAGGCAGA GGCCCACCAG CTCCAGCAGG CACCCTGACA GTCTGTCAT 2941 CTGTTGAGAAT GCTATGAAGT ATAACCCTGC ATATGAGAGC CACTTGGCTG GATGTGAAC 3001 TGTATGAGAAA TCCTATAGCC AACATCCCTT CTATAGCTCT GCTAGTGAAG AGGTGATTG 3061 GTGGTCTGAGC AGAGAAGAAA TCAGACAGAT GTATGAAAGT AGCGACCTTT CCAAAGAGG 3121 AAATTCAAGAG AGAATGAGGA TTTTGGAACT CTATGCTAAT GATCCTGAGT TTGCAGCTT 3181 TTGTGAGAGAG CATCAAATGG AGGAGCTTTA ACTTAAATGC CTGATTCTTG ACACCAATC 3241 AGAAGCTTGGA GAAGATGGAG AAGGCTTGTT CTCTCTGCTG TTTAACTAAT CCAGAAGAA 3301 GAGTTTGTATT GAAGAATAAA TAAGGAAACA TGGGACGCAC TTCTCATTCC AACACTGCA 3421 AGAGATGTGTC AAAAGAAAAT TAAAGTGGTG TGAACTCTGA TTTTGTAACA TATTCTAAA 3481 AGCAAACAAAT AAAACAGAAC CAAACCAAAA GCTTAAAGCC AGACCTTGGA GTTGGGGCT 3541 GCAGTGCCTCT GACTCTGACT TTTTGAGAGC ATCTCTAAGA ACTATGGCCC AGGCTTTCT 3601 AGTAAGAACAT AAAGTGAGAC TAATGAGTAA AGCTTAGAAT GCGACTGTTT TGTGACATA 3661 CTCGTTAAAGT CGAATGAGAT AGAGGAAGCT TTGAAGTAAT TTTAATATAG TTTAAACTC 3721 AAACACTCATC TAAATAAAAA TTAGGCTTTT GGAACAGATT GCTGAGTCAG GCAATCTTT

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3781 AGGTGCAGTAT ATCTTGTTTA TGTTTGATGC TTGCTTCCTA TCTGTTCTTG AGCTTCTTG
3841 AGCCCATAGAT CAAGACTACA ATGCTCTTAA ATTAGTTATG TCAATATTTG CCACAGTTT
3901 GGTCCTCAATT AGGCACCCTT AAGAGGAAGC AAATTGAGGA ATTNCNNTTC ATCAGCTTG
3961 GTTTGTGGACA TACCAGTGGG CCTTTTTCTT GATTATTAAT TGATGTAGAA AGGCCCAGC
4021 TCACTATGGGT GGTACTATCC TTAGGCAGGG GTTTGGGGAG TTAAGTTGCA AAAGAAAGG
4081 TAAAAGCCAGCT ACAAGAAGCC AGCCAATAAG CACTTTCCTT TGTGGTTTCT TCTTCAAAC
4141 TCCTGTCTTGG CTTCTCTCA TGGTAGACTA TAACCTATAA GCCAAATAAA CTCTTTCTT
4201 GGAA
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SEQ ID NO:24 Mouse IPM 200 amino acid sequence (partial)

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LOCUS MOUSE IPM 200 1069 AA PROT
DEFINITION Mus musculus IPM 200 core protein, lacks NH-terminus
KEYWORDS -
FEATURES From To Definition
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1 VANHVKYFKA RVCQEAIWEA FRTFWDRLPG RDEYRHWMNL CEDGVTSVFE MGAHFSQSVE
 61 HRNLIMKKLA YTREAESSSC KDQSCGPELS FPVPIGETST LTGAVSSASY PGLASESSAA
121 SPQESISNEI ENVTEEPTQP AAEQIAEFSI QLLGKRYSEE LRDPSSALYR LLVEEFISEV
181 EKAFTGLPGY KGIRVLEFRA PEENDSGIDV HYAVTFNGEA ISNTTWDLIS LHSNKVENHG
241 LVEMDDKPTA VYTISNFRDY IAETLHQNFL MGNSSLNPDP KPLQLINVRG VLLPQTEDIV
301 WNTQSSSLQV TTSSIXVLQP DLPVAPEGRT SGSFILEDGL ASTEELEDTS IDGLPSSPLI
361 QPVPKETVPP MEDSDTALLS TPHLTSSAIE DLTKDIGTPS GLESLASNIS DQLEVIPWFP
421 DTSVEKDFIF ESGLGSGSGK DVDVIDWPWS ETSLEKTTKP LSKSWSEEQD ALLPTEGREK
481 LHIDGRVDST EQIIESSEHR YGDRPIHFIE EXSHVRSTIP IFVESATPPT SPIFSKHTSD
541 VPDIDSYSLT KPPFLPVTIA IPASTKKTDE VLKEDMVHTE SSSHKELDSE VPVSRPDMQP
601 VWTMLPESDT VWTRTSSLGK LSRDTLASTP ESTDRLWLKA SMTQSTELPS TTHSTQLEEE
661 VIMAVQDISL ELDQVGTDYY QSELTEEQHG KADSYVEMST SVHYTEMPIV ALPTKGGVLS
721 HTQTAGALVV FFSLRVTNML FSEDLFNKNS LEYKALEQRF LELLAPYLQS NLSGFQNLEI
781 LSFRNGSIVV NSRVRFAESA PPNVNKAMYR ILEDFCTTAY QTMNLDIDKY SLDVESGDEA
841 NPCKFQACNE FSECLVNPWS GEAKCKCYPG YLSVDELPCQ SLCDLQPDFC LNDGKCDIMP
901 GHGAICRCRV GSNWWYRGQH CEEFVSEPFV IGITIASVVS FLLVASAVVF FLVKMLQAQN
961 VRRERQRPTS SSRHPDSLSS VENAMKYNPA YESHLAGCEL YEKSYSQHPF YSSASEEVIG
1021 GLSREEIROM YESSDLSKEE IQERMRILEL YANDPEFAAF VREHOMEEL
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SEQ ID NO:25 Monkey IPM 200 cDNA sequence (partial)

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05-0CT-1998
                                                     SYN
            MONKEY IPM
                          546 BP SS-DNA
LOCUS
           Macaca fascicularis IPM 200 cDNA, 5' end
DEFINITION
ACCESSION
KEYWORDS
                  From
                        Τo
                               Definition
FEATURES
                               IPM 200
                  188
                         546
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1 GAATTCGGCT TGGACAACCA TTTTTCTTTC CGCTAATTTA TAATGGTTTT GAAGTGGTTG
61 TTCATTCTCA AACATAGACT TTTAAATGTT AGGTCTTTCC TATAACTTGT TGTTATTGGA
121 AGTTTCAAGG ATTTGGACGC TCAGTTAAGG ATTTTGTCCT CTCCTCATTC CTTTGGTTTT
181 GGCCCAAATG ATTATGTTTC CTCTTTTTG GAAGATTTCT CTGGGTATTT TGATATTTGT
241 CCTGATAGGA GACTTTCCAT CGTTAACAGC ACAAACCTAC TTATCTTTAG AGGAGATCCA
301 AGAACCCAAG AGTGCAGTTT CTTTTCTCCT GCCTGAAGAA TCAACAGACC TTTCTCTAGC
361 TACCAAAAAG AAACAGCCTC TGGACCTCAG AGAAACTGAA AGACAGTGGT TACTCAGAAG
421 GCGGAGATCT ATTCTGTTTC CTAATGGAGT AAAAATCTGC CCAGATGAAA GTGTTACAGA
481 GGCTGTGGCA AATCATGTGA AGTATTTTAA AGTCCGAGTG TGTCAGGAAG CTGTCTGGGA
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11

SEQ ID NO:26 Monkey IPM 200 amino acid sequence (partial)

16-0CT-1998 PROT SYN MONKEY IPM 119 AA LOCUS DEFINITION Macaca fascicularis IPM 200 core protein, NH-terminal fragment ACCESSION KEYWORDS From To Definition **FEATURES** 114 Native NH-terminus DOMAIN 82

1 MIMFPLFGKI SLGILIFVLI GDFPSLTAQT YLSLEEIQEP KSAVSFLLPE ESTDLSLATK 61 KKQPLDLRET ERQWLLRRRR SILFPNGVKI CPDESVTEAV ANHVKYFKVR VCQEAVWEK

SEQ ID NO:27 Human IPM 150 isoform A variant cDNA sequence (3261 bp)

taaaccaaga aggttatcct caatcatctg gtatcaatat ataattattt ttcctttntg ttacttttta atgagatttg aggttgttct gtgattgtta tcagaattac catgcacaaa									60 120							
agcc	aga	atq	tat	tta	qaa	act	aga	aga	gct	att	ttt	gtt	ttt Phe	tgg	att	169
ttt Phe	Leu	caa Gln	Val	Gln	Gly 20	Thr	Lys	Asp	Ile	Ser 25	att Ile	Asn	Ile	Tyr	His 30	217
tct Ser	gaa Glu	act Thr	aaa Lys	gac Asp 35	ata Ile	gac Asp	aat Asn	ccc Pro	cca Pro 40	aga Arg	aat Asn	gaa Glu	aca Thr	act Thr 45	gaa Glu	265
agt Ser	act Thr	gaa Glu	aaa Lys 50	atg Met	tac Tyr	aaa Lys	atg Met	tca Ser 55	act Thr	atg Met	aga Arg	cga Arg	ata Ile 60	ttc Phe	gat Asp	313
ttg Leu	gca Ala	aag Lys 65	cat His	cga Arg	aca Thr	aaa Lys	aga Arg 70	tcc Ser	gca Ala	ttt Phe	ttc Phe	cca Pro 75	acg Thr	gjà aaa	gtt Val	361
aaa Lys	gtc Val 80	tat	cca Pro	cag Gln	gaa Glu	tcc Ser 85	atg Met	aaa Lys	cag Gln	att Ile	tta Leu 90	gac Asp	agt Ser	ctt Leu	caa Gln	409
gct Ala 95	tat	tat Tyr	aga Arg	ttg Leu	aga Arg 100	gtg	tgt Cys	cag Gln	gaa Glu	gca Ala 105	gta Val	tgg Trp	gaa Glu	gca Ala	tat Tyr 110	457
caa	atc Ile	ttt Phe	ctg Leu	gat Asp 115	cgc	atc Ile	cct Pro	gac Asp	aca Thr 120	gly aaa	gaa Glu	tat Tyr	cag Gln	gac Asp 125	tgg Trp	505
gtc Val	agc Ser	atc Ile	tgc Cys 130	cag Gln	cag Gln	gag Glu	acc Thr	ttc Phe 135	tgc Cys	ctc Leu	ttt Phe	gac Asp	att Ile 140	gga Gly	aaa Lys	553
aac Asn	ttc Phe	agc Ser 145	aat Asn	tcc Ser	cag Gln	gag Glu	cac His 150	ctg Leu	gat Asp	ctt Leu	ctc Leu	cag Gln 155	cag Gln	aga Arg	ata Ile	601
Lys	Gln 160	Arg	Ser	Phe	Pro	Asp 165	Arg	Lys	Asp	Glu	Ile 170	Ser	gca Ala	Glu	Lys	649
aca Thr 175	tta	gga Gly	gag Glu	cct Pro	ggt Gly 180	gaa Glu	acc Thr	att Ile	gtc Val	att Ile 185	tca Ser	aca Thr	gat Asp	gtt Val	gcc Ala 190	697

Asn	Val	Ser	Leu	Gly 195	Pro	Phe	Pro	Leu	Thr 200	Pro	Asp	gac Asp	Thr	Leu 205	Leu	745
Asn	Glu	Ile	Leu 210	Asp	Asn	Thr	Leu	Asn 215	Asp	Thr	Lys	atg Met	Pro 220	Thr	Thr	793
gaa Glu	aga Arg	gaa Glu 225	aca Thr	gaa Glu	ttc Phe	gct Ala	gtg Val 230	ttg Leu	gag Glu	gag Glu	cag Gln	agg Arg 235	gtg Val	gag Glu	ctc Leu	841
agc Ser	gtc Val 240	tct	ctg Leu	gta Val	aac Asn	cag Gln 245	aag Lys	ttc Phe	aag Lys	gca Ala	gag Glu 250	ctc Leu	gct Ala	gac Asp	tcc Ser	889
cag Gln 255	tcc	cca Pro	tat Tyr	tac Tyr	cag Gln 260	gag Glu	cta Leu	gca Ala	gga Gly	aag Lys 265	tcc Ser	caa Gln	ctt Leu	cag Gln	atg Met 270	937
caa	aag Lys	ata Ile	ttt Phe	aag Lys 275	aaa	ctt Leu	cca Pro	gga Gly	ttc Phe 280	aaa Lys	aaa Lys	atc Ile	cat His	gtg Val 285	tta Leu	985
gga Gly	ttt Phe	aga Arg	cca Pro 290	aaq	aaa Lys	gaa Glu	aaa Lys	gat Asp 295	ggc Gly	tca Ser	agc Ser	tcc Ser	aca Thr 300	gag Glu	atg Met	1033
caa Gln	ctt Leu	acg Thr 305	qcc	atc Ile	ttt Phe	aag Lys	aga Arg 310	cac His	agt Ser	gca Ala	gaa Glu	gca Ala 315	aaa Lys	agc Ser	cct Pro	1081
gca Ala	agt Ser 320	qac	ctc Leu	ctg Leu	tct Ser	ttt Phe 325	gat Asp	tcc Ser	aac Asn	aaa Lys	att Ile 330	gaa Glu	agt Ser	gag Glu	gaa Glu	1129
gtc Val 335	tat	cat His	gga Gly	acc Thr	atg Met 340	gag	gag Glu	gac Asp	aag Lys	caa Gln 345	cca Pro	gaa Glu	atc Ile	tat Tyr	ctc Leu 350	1177
aca	gct Ala	aca Thr	gac Asp	ctc Leu 355	aaa	agg Arg	ctg Leu	atc Ile	agc Ser 360	aaa Lys	gca Ala	cta Leu	gag Glu	gaa Glu 365	gaa Glu	1225
caa Gln	tct Ser	ttg Leu	gat Asp 370	gtg	gjà aaa	aca Thr	att Ile	cag Gln 375	ttc Phe	act Thr	gat Asp	gaa Glu	att Ile 380	gct Ala	gga Gly	1273
tca Ser	ctg Leu	cca Pro 385	gcc Ala	ttt Phe	ggt Gly	cct Pro	gac Asp 390	acc Thr	caa Gln	tca Ser	gag Glu	ctg Leu 395	ccc Pro	aca Thr	tct Ser	1321
ttt Phe	gct Ala 400	gtt Val	ata Ile	aca Thr	gag Glu	gat Asp 405	gct Ala	act Thr	ttg Leu	agt Ser	cca Pro 410	gaa Glu	ctt Leu	cct Pro	cct Pro	1369
Val 415	gaa Glu	Pro	Gln	Leu	Glu 420	Thr	Val	Asp	Gly	Ala 425	Glu		Gly	Leu	Pro 430	1417
gac	act	tct Ser	tgg Trp	tct Ser 435	Pro	cct Pro	gct Ala	atg Met	gcc Ala 440	Ser	acc Thr	tcc Ser	ctg Leu	tca Ser 445	gaa Glu	1465
Āla	Pro	Pro	Phe 450	Phe	Met	Ala	Ser	Ser 455	Ile	Phe	Ser	Leu	Thr 460	Asp	caa Gln	1513
Gly	Thr	Thr 465	Asp	Thr	Met	Ala	Thr 470	Asp	Gln	Thr	Met	Leu 475	. Val	Pro	ggg	1561
ctc Leu	acc Thr 480	atc Ile	ccc	acc Thr	agt Ser	gat Asp 485	Tyr	tct Ser	gca Ala	ato Ile	ago Ser 490	Gln	ctg Leu	gct Ala	ctg Leu	1609

J _	-	Ser Ser Asp	gac agc cga tca agt gca Asp Ser Arg Ser Ser Ala 505 510	1657						
Gly Gly Glu Asp	Met Val Arg 515	His Leu Asp (gaa atg gat ctg tct gac Glu Met Asp Leu Ser Asp 525	1705						
Thr Pro Ala Pro	Ser Glu Val	Pro Glu Leu S 535	agc gaa tat gtt tct gtc Ser Glu Tyr Val Ser Val 540	1753						
			gtc tca gct tta cag tat Val Ser Ala Leu Gln Tyr 555	1801						
	_	_	aag ggc cga gag ctg gta Lys Gly Arg Glu Leu Val 570	1849						
J J		Ala Asn Met A	gcc ttc tcc aac gac ctg Ala Phe Ser Asn Asp Leu 585 590	1897						
			etg gag caa caa ttc aca Leu Glu Gln Gln Phe Thr 605	1945						
	Pro Tyr Leu		ett aca gga ttt aag caa Leu Thr Gly Phe Lys Gln 620	1993						
			gtg att gtg aat agc aaa /al Ile Val Asn Ser Lys 635	2041						
		_	etc acc aag gct gtg cac Leu Thr Lys Ala Val His 650	2089						
		Ser Ala Ala A	gcc caa caa ctc cat ctg Ala Gln Gln Leu His Leu 665 670	2137						
		_	cca gct gat caa gca gat Pro Ala Asp Gln Ala Asp 685	2185						
• •			gcc caa tgt gta aag aac Ala Gln Cys Val Lys Asn 700	2233						
			aaa cca gga tat gac agc Lys Pro Gly Tyr Asp Ser 715	2281						
			etc tgt ggc ctg gca caa Leu Cys Gly Leu Ala Gln 730	2329						
		Glu Arg Glu I	etc cat gcg gtt cca gat Leu His Ala Val Pro Asp 745 750	2377						
	_		gtt aaa agt tcc aaa atc /al Lys Ser Ser Lys Ile 765	2425						
aac aaa ata aca Asn Lys Ile Thr 770		taa aagaaattct	gaattactga ccgtagaata	2480						
tgaagaattt aaccatcaag attgggaagg aaattaaaaa ctgaaaatgt acaattatca 254										
cttaggctat ctcaagagag atgatttgcc ttctcaagga aaatggagac aggcatattc 2										
atgggtcatc aaaatccaga catacagtca acactgagaa tcagcacaca ccatatttca 2660										

aatatagaag agtcatgtac ttggcaacca gtaaattctg aaaaaaaaga cacttactta 2720 ttattaaaac cccaaatgca atcagcgaaa catatttta ctattcttgg atgatagtca 2780 aaatgatcat aagccaggtt tgcttccacc ttccctgaaa atttactca cagatcattt 2840 gcaacaagca tagcttactt attgtttagg gactgaacaa tttattggga agcaaactct 2900 ttatatgcta gaaagtacat ttaaaagatg actacttacg cagggagatg caggtctctc 2960 taaacgcatg aatgtatgta gtgtgtaggc actgtagtga gtgtatatat gctccacact 3020 acgtctgata aacacaaacc tcagtattca gttattaggc acactagttt tatacgcaac 3080 tactgcttac atagtagact gttttgttgc caataatctt tgaattgttc tttaaaagaa 3140 actgaggttc agatacacat accatggaaa aatcttactt ttcttgttac tacacaaagc 3200 tattttaaag aagatgctat gttgggagaa gggcgaagtt gtactatatg acataatcaa 3260 t

SEO ID NO:28 Human IPM 150 isoform A variant amino acid sequence

Met Tyr Leu Glu Thr Arg Arg Ala Ile Phe Val Phe Trp Ile Phe Leu Gln Val Gln Gly Thr Lys Asp Ile Ser Ile Asn Ile Tyr His Ser Glu Thr Lys Asp Ile Asp Asn Pro Pro Arg Asn Glu Thr Thr Glu Ser Thr Glu Lys Met Tyr Lys Met Ser Thr Met Arg Arg Ile Phe Asp Leu Ala Lys His Arg Thr Lys Arg Ser Ala Phe Phe Pro Thr Gly Val Lys Val Cys Pro Gln Glu Ser Met Lys Gln Ile Leu Asp Ser Leu Gln Ala Tyr Tyr Arg Leu Arg Val Cys Gln Glu Ala Val Trp Glu Ala Tyr Arg Ile Phe Leu Asp Arg Ile Pro Asp Thr Gly Glu Tyr Gln Asp Trp Val Ser Ile Cys Gln Gln Glu Thr Phe Cys Leu Phe Asp Ile Gly Lys Asn Phe Ser Asn Ser Gln Glu His Leu Asp Leu Leu Gln Gln Arg Ile Lys Gln Arg Ser Phe Pro Asp Arg Lys Asp Glu Ile Ser Ala Glu Lys Thr Leu Gly Glu Pro Gly Glu Thr Ile Val Ile Ser Thr Asp Val Ala Asn Val Ser Leu Gly Pro Phe Pro Leu Thr Pro Asp Asp Thr Leu Leu Asn Glu Ile Leu Asp Asn Thr Leu Asn Asp Thr Lys Met Pro Thr Thr Glu Arg Glu Thr Glu Phe Ala Val Leu Glu Glu Gln Arg Val Glu Leu Ser Val Ser Leu Val Asn Gln Lys Phe Lys Ala Glu Leu Ala Asp Ser Gln Ser Pro Tyr Tyr Gln Glu Leu Ala Gly Lys Ser Gln Leu Gln Met Gln Lys Ile Phe Lys Lys Leu Pro Gly Phe Lys Lys Ile His Val Leu Gly Phe Arg Pro Lys Lys Glu Lys Asp Gly Ser Ser Ser Thr Glu Met Gln Leu Thr Ala Ile Phe Lys Arg His Ser Ala Glu Ala Lys Ser Pro Ala Ser Asp Leu Leu Ser Phe Asp Ser Asn Lys Ile Glu Ser Glu Glu Val Tyr

His Gly Thr Met Glu Glu Asp Lys Gln Pro Glu Ile Tyr Leu Thr Ala Thr Asp Leu Lys Arg Leu Ile Ser Lys Ala Leu Glu Glu Glu Gln Ser Leu Asp Val Gly Thr Ile Gln Phe Thr Asp Glu Ile Ala Gly Ser Leu Pro Ala Phe Gly Pro Asp Thr Gln Ser Glu Leu Pro Thr Ser Phe Ala Val Ile Thr Glu Asp Ala Thr Leu Ser Pro Glu Leu Pro Pro Val Glu Pro Gln Leu Glu Thr Val Asp Gly Ala Glu His Gly Leu Pro Asp Thr Ser Trp Ser Pro Pro Ala Met Ala Ser Thr Ser Leu Ser Glu Ala Pro Pro Phe Phe Met Ala Ser Ser Ile Phe Ser Leu Thr Asp Gln Gly Thr Thr Asp Thr Met Ala Thr Asp Gln Thr Met Leu Val Pro Gly Leu Thr Ile Pro Thr Ser Asp Tyr Ser Ala Ile Ser Gln Leu Ala Leu Gly Ile Ser His Pro Pro Ala Ser Ser Asp Asp Ser Arg Ser Ser Ala Gly Gly Glu Asp Met Val Arg His Leu Asp Glu Met Asp Leu Ser Asp Thr Pro Ala Pro Ser Glu Val Pro Glu Leu Ser Glu Tyr Val Ser Val Pro Asp His Phe Leu Glu Asp Thr Thr Pro Val Ser Ala Leu Gln Tyr Ile Thr Thr Ser Ser Met Thr Ile Ala Pro Lys Gly Arg Glu Leu Val Val Phe Phe Ser Leu Arg Val Ala Asn Met Ala Phe Ser Asn Asp Leu Phe Asn Lys Ser Ser Leu Glu Tyr Arg Ala Leu Glu Gln Gln Phe Thr Gln Leu Leu Val Pro Tyr Leu Arg Ser Asn Leu Thr Gly Phe Lys Gln Leu Glu Ile Leu Asn Phe Arg Asn Gly Ser Val Ile Val Asn Ser Lys Met Lys Phe Ala Lys Ser Val Pro Tyr Asn Leu Thr Lys Ala Val His Gly Val Leu Glu Asp Phe Arg Ser Ala Ala Ala Gln Gln Leu His Leu Glu Ile Asp Ser Tyr Ser Leu Asn Ile Glu Pro Ala Asp Gln Ala Asp Pro Cys Lys Phe Leu Ala Cys Gly Glu Phe Ala Gln Cys Val Lys Asn Glu Arg Thr Glu Glu Ala Glu Cys Arg Cys Lys Pro Gly Tyr Asp Ser Gln Gly Ser Leu Asp Gly Leu Glu Pro Gly Leu Cys Gly Leu Ala Gln Arg Asn Ala Arg Ser Ser Arg Glu Arg Glu Leu His Ala Val Pro Asp His Ser Glu Asn Gln Ala Tyr Lys Thr Ser Val Lys Ser Ser Lys Ile Asn Lys Ile Thr Arg